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Sequence 31vl BP; 857 A; 531 C; 451 G; 1222 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page my-bu; 95pp; Englist.
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freat tancer protein useful in the detection and treatment of
cancer
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02-JUL-1999; 990S-0346327
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breast canber; vaccine; ss.
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    score 373; Lb 21; Length 3101;
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actived from the 5' ends of maths and even in those cases where longer cDAN sequences have been obtained, the full 5' UTR is rarely included.
5' ESTS are derived from makes with thotat 5' ends and can therefore be used to obtain full length cDNAs and genome DNAs. 5' ESTS are also used to obtain full length cDNAs and genome to DNAs. 5' ESTS are also used to distance the sequence of the sequences and the sequences and to design expression and secretion vectors.

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24-0CT-2001 (first entry)

AAS10873 standard:

CDNA; 41100 BP

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Query Match 3.2%; Sc Best Local Similarity 100.0%; P Matches 32; Conservative 0;

Score 32: DB 21; Pred. No. 0.0013; 9: Mismatches 0

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Length 305:

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The present sequence is one of a large number of 5' ESTS derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTS were prepared from total human RNAs or polyA- RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from olivo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-SEP-2000
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                                                                                                                                                                                                                  Claim 1; SEQ ID 31332; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                            New nucleic acid that is a 5' expressed sequence tag (5' ESI) for obtaining cDNAs and genomic LNAs that currespond to 5'ESIs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-500381/45
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gene therapy; chromosome mapping; ss.
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AAC27257 standard;
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12-JAN-2000
13-JAN-2000
07-AUG-2000
04-JAN-2001
The invention relates to nucleic acids encoding NOVX (k being an integer from 1-8) polypeptides. The NOVX nucleic acids and polypeptides are useful in diagnosing, treating or manufacturing a medicament for a dissase or disorder associated with NOVX expected as medicant no (cancer and diabetic returopathy), anglogenic or pulmonary disorders, fertility
                                                                                                               Nucleic acids encoding polypeptides, designated NOVX polypeptides, useful for treating a syndrome associated with a NOVX-associated discrete, e.g. cell proliferation (e.g. cancer and disbetic retinopathy), angiogenic or pulmonary disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human: NOV6: ds: 78347267; fertility disorder: spermatogenesis; cardiant: cytostatic; immunomobulatory; entiproliferative; antidiabetic; cell proliferation; cencer; diabetic retinopathy; angiogenic disorder; pulmonary disorder; nacmotopietic disorder; immunological disorder; inflammatory disorder; tumour related disorder; emphysema; cirricosis; wound healing; gene therapy.
                                                                                    Disclosure: Page 22-30; 144pp; English.
                                                                                                                                                                                                                  Prayaga SK.
MacDougall J:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genomic DNA containing the NOV6 gene.
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disorders (e.g. of spermatoyenesis), hematopoietic, imminisperia, citrin.sis, which inflammatory and tumbur related disorders, emphysema, citrin.sis, which healing, NOVA nucleic acids are also useful in gene therapy. They are also useful for screening for a modulator of activity or interapy. They are also useful for determining the presence of or predisposition to a NOVA-associated disorder. The present seyuence contains the year for NOV6 (78847267), which has sequence homology to interleukin a.

Sequence 41100 BP; 12250 A; 8060 C; 8159 G; 12631 T; 0 other;

Query Ma Best Loc Matches

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Score 30; DB; Pred. No. 0.0

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Movel human secreted proteins useful for diagnosis, prevantion treatment of disorders including neurological, cell proliferaticardiovascular, autoimmune/inflammatory disorders and microbial infections
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03-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 362; 427pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-MAR-2000; 2000WO-US06059
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted protein cDNA sequence #45.
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DB; AAB27724.
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99US-0168622.
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Best Local S
Matches 29
                                                                                                                                                                             thatrix. Liest, pastrointestinal tract, liver, lung, or urogenital;
(b) immine disorders e.g. Addison's disease, ellergies, autoimmune
haemolytta ansemia, autoimmune thyroiditis, diabetes mellitus, Crohn's
disease, multiple sclerosis, rieumatoid arthritis and ulcerative
colitis; (c) caratovascular disorders such as myocatdal ischaemias; (d)
"Journ healing; (e) resurological diseases e.g. cerebral anoxia and
epileysy; and (f) infectious diseases such as viral, bacterial, fungal
and parasitic infections.
Sequence 1308 BP; 322 A; 227 C; 286 G; 472 T; 1 other;
                                                               on 2.9%; So 1 Similarity 100.0%; F
                                                             Score 29; DB; Pred. No. 0.0
0; Mismatches
                                                                              DB 21;
0.017;
                                                                   Ç,
                                                                                              Length 1308;
                                                                 Indels
                                                                 0;
                                                               Gaps
                                                                 0;
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Human train Expressed Sequence Tag EST01493 16-MAK-1994 AAQ59633 standara; CDNA; 315 BF

Homo sapiers #09316178-A

Gene transcription product; genetic markers; tagging; in vivo; transcription; mapping; locations; chromosomes; chromosomal; ss

19-400-1993

12-FEB-1993; 93W0-US01294

12-FEB-1992; 92US-0837195

(USSH ) US DEPT HEALTH & HUMAN SERVICE

Adams MD, Moreno RF, Venter CJ;

Enricia digonuclectides and corresp, sequences - used markers for human genes transcribed in vivo, facilitate of most human genes

as tagging

Example 4; Page 207; 500pp; English.

The Expressed Sequence Tag was isolated from a human brain cDNA library as part of a large set of ESIS which can be used as markers for human genes transcribed in vivo. They can be used to facilitate tagging of most human genes, for mapping locations of expressed genes can chromosomes, for individual or forensic identification, for mapping locations of disease associated genes, for identification of tissue type, and for preph. of antisense sequences, probes and constructs. ESTU153 has a "poor" coding probability as evaluated using the coding-region prediction program CRM. See also AAQS9041-061440.

Sequence 319 BP; 118 A; 42 C; 75 G; 62 T; 2 other;

Ouery Match 2.7%; So Best Local Similarity 100.0%; i Matches 27; Conservative 0; Score 27; DB 14; Pred. No. 0.12; 0; Mismatches 0 0; Length 319; Indels 0 Gaps

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RESULT

AAF92122

AAF92122:

XX AAF92122:

XX AAF92122:

XX AAF92122:

XX Human PRO1928 cDHA.

XX PR Human PRO1928 cDHA.

XX Human PRO1928 cDHA.

XX PR Human PRO1928 cDHA.

XX PR Human PRO1928 cDHA.

XX DOS-MAR-2001.

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XX DOS-MAR-2001.

XX CETH JORD JORDON-USCHILL

PR 11-184-2001.

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                                                              The present sequence is the coding sequence for a human PRO polypertial (secreted and transmembrane). The PRO protein, and PRO agonists, FRO antibodies are useful for preparation of a medicament useful in the treatment of a condition which is responsive to protein agonists, antagonists or anti-PRO antibodies. The PRO protein may also be employed as molecular weight markers for protein electrophoresis. The PRO coding sequence has applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping.
                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Fig 129; 278pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Eighty four nucleic acids encoding PRO polypeptides ...setul molecular biology, including use as hybrivization proces, archromosome and gene mapping.
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, Watanabe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8
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            I; 0 other:
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Query Match 2.7%: Best Local Similarity 100.0%; Matches 27; Conservative

t; Score 27; bb t; Pred. No. 0.1 0; Mismatches

22; Length 1177; ن. Indels

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AADU5579-AADU5556 represent chiks corresponding to 28 human secreted protein genes and AADU1770-AADU1849 represent the proteins they encourable protein genes and AADU1770-AADU1849 represent the proteins they encourable proteins are useful for preventing. The genes and their secreted proteins are useful for preventing, treating and tail granting medical conditions, e.g., by protein or generating perfecting and tail conditions, e.g., by protein or generating perfection in a sample or by determining the presence of mutations in the new genes, packific uses are described for each of the about to the new genes, packific uses are described for each of the about the new genes, the presence of the pre

Claim 1; Page 474; 567pp; English.

Kwe nubjets acht melecules encoding numen secteted proteins, used preventing, treating or nmeliorating a disorder, e.g. Alzheimer's Parkinson's diseases and cancers'

12-ROV-1955; 550S-0164744. 36-JUN-2006; 2660US-6215140. 08-NUV-2000; 2000WO-US30628

(HUMA-) HUMAN GENUME SCI INC

Ruben SM.

Komatsoulis GA,

Baker KF,

Young r

PSLB;

2301-316491/33. 55: AAE01838.

17-MAY-20ul

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Human: secreted protein: prollierative disorder: cancer: tumour: foetal accormality; developmental accormality; hacmatopoletic disorder: manne system disorder: AIDS; autoimmune disease; rheumatoid arthritis; inflammation; allergy; neurological disorder: Alzhelmer's disease; parkinsor: alsease; cognitive disorder: solizophenala astima; skin disorder: psoriasis; sepsis; dishetes; atheroselerosis; cardicohacular disorder: sangiogenic disorder; kinney disorder: pastrointestinal disorder: pregnancy related disorder: pastrointestinal disorder; pregnancy related disorder: pastrointestinal disorder; pregnancy related disorder; call culture; chemotaxis; food additive; gene therapy; call culture; chemotaxis; food additive; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted protein-encoding gene 21 cDNA clone HDPTN65, SEQ ID NO:75
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/product- "Human
                                                                                                                                                                                                                                                                                                                           mature secreted protein.
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Human; secreted protein; proliferative disorder; cancer; tumour; foetal abnormality; developmental abnormality; haematopoiette ulisorder; immune system disorder; AIDS; autoimmune disease; rheumatoid arthritis; inflammation; allergy; neurological disorder; Althelmer's disease; parkinson's disease; cognitive disorder; schizophrenia; astlang; skin disorder; psoriasis; sepsis; diabetes; atherosclerosis; cardiovascular disorder; angiogenic disorder; kidney disorder; gastrointestinal disorder; prephancy related disorder; gene therapy; endocrine disorder; infection; wound healing; vulnerary; cell culture; chemotaxis; food additive; binding partner identification; ss.
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12-NOV-1999; 990S-0164744
30-JUN-2000; 20000S-0215140
                                                                                                                             17-MAY-2001.
                                                                                                                                                                                WO200134627-A1
                                                                                                                                                                                                                                                                                                                               sig_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted protein-encoding gene 21 cDNA clone HDPTW65, SEQ ID NO:31
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/product * Human secreted protein*
136.249 b
261.379
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/product= "Human mature secreted protein"
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ce abnormalities, haumatopoiette disorders, diseases of the interme system, AIDS, autoimmune diseases (w.g., rhematoid arthritis), initialmation. AIDS, autoimmune diseases (w.g., rhematoid arthritis), initialmation. Ce alteries, neurological disorders (e.g., Milheimar's diseases, established as schizohrenia, astinna, parkinson's disease), comitive disorders, schizohrenia, astinna, ce skin disorders (e.g., psorialsis), sepsis, diabetes, atheroscierosis, cardiovascular disorders, angiogenic disorders, atheroscierosis, cardiovascular disorders, angiogenic disorders, atheroscierosis, compatent standard and compatent disorders, and infections. The proteins can also be used to alid wound chealing and epithelial cell proliferation, to prevent skin aging due to culture of primary tissues, to regenerate tissues, to deprify their compate ligands or binding partners, and in chemotaxis, and can be used to as a food additive or preservative to monify storage properties.

Cantibodies specific for a protein of the invention can be used in diagnostic immunosorbent assay (ELISA). The present sequence represents a human considerated with the disorders mentioned above, and cinamonosorbent assay (ELISA). The present sequence represents a human exception of the invention can be used to secreted protein-encodiny cDNA of the invention.
Claim 1; Fage 436-437; 567pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New runcleic acid molecules encoding human secreted proteins, used preventing, treating or ameliorating a disorder, e.g. Alzheimer's Farkinson's diseases and cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI: 2001-316451/33.
P-PSDB: AAE01750.
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Query Match 2.7%; Sc Best Local Similarity 100.0%; P Matches 27; Conservative 0;

Score 27; 16 22; Length 1102; Pred. No. 0:11; 0; Mismatches 0; Indels

0; Indels v,

Sequence 1182 BP; 274 A; 170 C; 209 G; 519 T; 10 other;

CC ALGO5579-AADD5656 represent cDNAs corresponding to 28 human secreted protein genes and AADD170-AADD1867 represent the protein they encode. AADD1866-AADD1866 represent human secreted protein fragments of variants. The genes and their secreted proteins are useful for preventing. Creating or ameliorating medical conditions, e.g., by protein or gene therapy. Pathological conditions can be diagnosed by determining the abount of the new genes. Specific uses are described for each of the CC amutations in the new genes. Specific uses are described for each of the CC and include developing products for the diagnosed by the presence of proliferative disorders, cancer, tumours, foctal and developmental proliferative disorders (e.g., rheunatoid arthitis), inflammation, albey and proteins are most highly expressed. Albey autonomic diseases (e.g., rheunatoid arthitis), inflammation, albey and a secondary disorders, callergies, neurological disorders (e.g., rheunatoid arthitis), inflammation, albey as a neurological disorders, sepsis, disbetes, artherosclerosis, callergies, and infections. The proteins can also be used to said wound constitute this orders, pregnancy related disorders, endowed the subdurf, to maintain organs before transplantation, for supporting due to subdurf, to maintain organs before transplantation, for supporting due to contain a protein contain subsective the subdurf of the invention can be used in an indicating symptoms associated vith the disorders mentioned above, and contained subvected vith the disorders mentioned above, and contained subvected vith the disorders mentioned above, and contained subvected vith the disorders mentioned above, and contained contained

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Sequence 1917 BP; 558 A; 286 C; 349 G; 724 T; 0 other;
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DB 22; Length 1917; . 0.1;

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AAH73960 standard; cDNA; 2781 BP.
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AAH73960;
XX
AC BHuman halix-destabilising enzyme C2
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AC Human, helix-destabilising enzyme C2
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AC Human, helix-destabilising enzyme C3
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AC Human, helix-destabilising enzyme C4
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AC Human, helix-destabilising enzyme C5
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AC Human, helix-destabilising enzyme C5
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AC Human, helix-destabilising enzyme C7
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AC Human, selix-destabilising enzyme C7
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AC Human, helix-destabilising enzyme C8
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AC Human, helix-destabilising enzyme C8
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AC Human, helix-destabilising enzyme 
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Human helix-destabilising enzyme C20 coding sequence.

Human; helix-destabilising enzyme C20; cytostatic; virucide; immunodulatory; antiinflammatory; haemostatic; gene therapy; malignant tumour; haemopathy; HIV infection; immunological disease; inflammation; ss.

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Page 8
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Best Local Similarity 100.
Matches 27; Conservative
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                                                                                 exon
                                                                                                                                                                  exon
                                                                                                                                                                                                                                                                         Dectin 7, dendritic cell; c-type lectin, modae; immunity; adjuvant; allergy; autoimmune disease; gene therapy; vaccine; diagnosis; drug screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2781 BP; 711 A; 637 C; 595 G; 837 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is the coding sequence for human helix-destabilising enzyme C20. The enzyme and its coding sequence are useful in the diagnosts and treatment of mallyanat tumour, haemopathy, HIV infection, immunological diseases and various inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human helix-destabilizing enzyme C20 and encoded polynuclectide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-483253/52.
P-PSDB; AAG64390.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BIOD-) BIODOOR GENE TECHNOLOGY LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JAN-2000; 2000CN-0111591.
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exon
                                                                                                                                                                                                                                                                                                                                  Mouse dectin-2 gene.
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                                                                                                                                                                                                                                                                                                                                                                                        AAV42558;
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633..727
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/number= 2
728..2223
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/number-1
[72..632
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147..171
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:: Pred. No. 0.098:

0: Mismatches 0: Indels
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Ariizemi K. Takashima A;
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P-PSDB; AAW63010.
                                                                                                                         (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                               20-DEC-1996;
                                                                                                                                     22-LEC-1997;
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                                                                                                                                                  ₩Ū$826332-AZ
                                                                                                                                96US-0772440
                                                                                                                                      97WO-US23761
                                                                                                                                                        /number* 3
2325..5746
/*tag= 9
/number* 3
5747..5698
/*tag* h
/number* 4
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and it acid encoding dendritic cell specific peptide(s) dectin-1 and it was unless to regulate immune response, as vaccine adjuvants, for diagnosis and drug screening

Claim 7; Page 159-165; 200pp; English.

Sequence 10409 BP; 2816 A; 1968 C; 1976 G; 3541 T; 108 other;

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RESULT 13
AAX63005
ID AAX630
XX
AC AAX630
XX
                                                                                                        Guery Match. 2.7%; Score 27; Lis 19; Length 10409; Best Local Similarity 100.0%; Bred, Mc. G.084; Matches 27; Conservative 0; Mismatches 0; Indels 0
             AAX63005;
                                       AAX83005 standard; DNA; 29604 BP
                                                                                                                                                                 0; Indels 0;
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RESULT 14
AAZ61532/c
IX AAZ61532/c
XX AAZ615
AX AAZ615
AX I9-JUN
XX UOXyloX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-APR-1996;
29-DEC-1995;
29-DEC-1995;
30-JAN-1996;
30-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence represents a fragment of the genomic sequence containing the coding region for the mouse KRN gene (AAX83001). The corresponding human gene (AAX83001) encodes a protein related to Werner's Symdiome. The products can be used for the detection and treatment of Werner's syndrome (M3), an autosmal recessive disorder with a complex phenotype, as well as related diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Fig 7; 153pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9724435-A1
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                                                                                                                             Oxytoxin expressed sequence tag; 5'-OT EST; obesity; fertility; male; transgenic animal; human late onset obesity; late onset viscaral obesity; male infertility; wasting; anorexia; cachexia; malabsorptive state; catabolic state; inflammatory condition; Crohn's disease; Al.S wasting; burn; cancer; bone disease; vasopressin; oxytocin; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-363671/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fu Y, Mulligan J, Oshima J, Schellenberg GD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DARW-) DARWIN MOLECULAR CORP
(OSHI/) OSHIMA J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 29604 BP; 7634 A; 5861 C; 5985 G; 10123 T; 1 other:
                               W0200005686 · A1
                                                                                Rattus sp.
                                                                                                                                                                                                                                                                                         Cosmid CVO14 containing rat vasopressin and oxytocin locus
                                                                                                                                                                                                                                                                                                                                          19-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                              AAZ61522;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ61522 standard; DNA; 44576 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid molecule encoding the WRN gene product useful for detection and treatment of Merner's syndrome, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 960S-0632175.
950S-0009409.
950S-0580539.
960S-0010835.
960S-0594242.
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AAI16817
ID AAI168:
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Best Local Similarity 100.0%; P
Matches 27; Conservative 0;
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Score 27: Db 21: Length 44576; Pred. NJ. 0.072; 0; Mismatches 0; Indels ^.

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Gaps

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04-FEB-2000; 20000S-0180312. 26-HAY-2000; 2000S-0207156. 30-JUN-2000; 2000S-0608408. 30-JUN-2000; 2000US-0632366. 03-MGC-2060; 2000US-0234687. 21-SEP-2000; 2000US-0236359.

30-JAN-2001; 2001WO-US00670

09-AUG-2001. W0200157276-A2 Homo sapiens. Probe: fullan: microarray: gene expression: cervical epithelial cell; cervical cancer; ss.

Probe +6750 for gene expression analysis in human cervical cell sample.

12-001-2001

(first entry)

AAI16817 standard; DNA; 535 BP

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12-AUG-1555;
06-MAY-1555;
                                                                                                                                                                          12-AUG-1999;
                                                                                                                                                                                    24-FEB-2000.
                                                                                                Claim 5; Page 129-154; 162pp; English.
                                                                                                          New anti-obesity polypeptide useful for treating obesity or infertility in maximals \cdot
                                                                                                                          WPI: 2000-224331/19.
                                                                                                                                     Robinson ICAF, Stoye JP, Flavell D,
                                                                                                                                               (MEDI-) MEDICAL RES COUNCIL.
Sequence 44576 BP; 12157 A; 10993 C; 10857 G; 10569 T; 0 other;
                                                                                                                                                           98GB-0017566
99GB-0010522
                                                                                                                                                                           95WO-GB02658
                                                                                                                                     Wells SE, Le Tissier
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04-OCT-2000; 2000GB-0024263

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Best Local S
Matches 25
                                                                                                  04-FEB-2000: 2000US-0180312
26-MAY-2000: 2000US-0207456
30-JUN-2000: 2000US-0608408
03-MAG-2000: 2000US-0512356
21-SEP-2000: 2000US-021659
27-SEP-2000: 2000US-021659
04-CCT-2000: 2000US-0044233.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to human single exon humble acid probes (SEMP). The present sequence is one such probe. The SEMPs are derived from human hetac cells. The SEMPs can be used to produce a single exon microarray, which can be used for measuring human gare expression in a sample derived from human cervical epithelial cells. By measuring yene expression, the probes are therefore useful in grading and/or steping of diseases of the cervix, notably cervical cancer.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from Wifel at fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                        probe; microarray: human; placenta; antenatal diagnosis;
genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 25; SEQ ID No 6750; Amapp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human genome-derived single exon nucleic acid probes useful for
analyzing gene expression in human cervical epithelial cells -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 535 BP; 144 A; 127 C; 70 G; 154 T; 0 other;
                                                                                                                                                                                                                                                                        09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                        Probe #9116 used to measure gene expression in human placenta sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAI40430;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAI40430 standard;
                                                                                                                                                                                                                                                                                                        WO200157272-A2
                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-488897/53.
                                    Penn SG,
                                                                    (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                      30-JAN-2001; 2001WO-US00663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            at Similarity 100.0%;
25; Conservative (
                                    Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                      Chen W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8
                                         Rank
                                         DR;
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0.72:
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Best Local S
Matches 25
                                                                                                                                           The present in mention relates to single exch nucleic acid probes (SENF). The present sequence is one such probe. The probes are useful for producting a microarray for predicting, measuring and displaying gene expression it, samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders.
                                                                                                                                                                                                                             Claim
                                                                                                                    Sequence 535 6P; 144 A; 127 C; 70 G; 194 T; 0 other;
2.5%;
| Similarity 100.0%;
| 25; Conservative
                                                                                                                                                                                                                             ID No 9116; 654pp; English
                                                      0
                                                    Score 25; DB; pred. No. 0.7; Mismatches
                                                                   DB 22; Length 535; 0.72;
                                                          0;
                                                          Indels
                                                           0
                                                          Gaps
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Probe: human: microarray; gene expression; cervical epithelial cell; cervical cancer; ss. Probe #6466 for gene expression analysis in human cervical cell sample. 30-JAN-2601; 20v1WO-US00670 05-406-2001. Homo Sapiens. WJ2U0157278-AŽ.

AAI16533 standard: DNA; 554 BP

12-001-2001 (first entry)

AAI16533;

04 FEB -2000: 260003-0180312 26-MKT -2000: 260003-0207456 30 -JUN-2000: 260003-0608408 03 -MG-2000: 260003-0623286 21 -SEP -2000: 260003-0234687 27 -SEP -2000: 260003-0234687 04 -CCT -2000: 260003-0234635 (MOLE-) HOLECULAR DYNAMICS INC

WPI: 2001-468501/53. renn so, Hatizel LK, Chen W. Kank Lik;

Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human cervical epithelial cells -

Claim 25; SEQ ID No 6466; 487pp; English.

The present invention relates to human single exon nucleic acid probes (SEMP). The present sequence is one such probes. The SEMPs are derived from human Hela cells. The SEMPs can be used to produce a single exon nicroarray, which can be used for measuring human gene expression in a sample defived from human cervical epithelial cells. By measuring pane expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIFO at tip.wipo.int/pub/published\_pct\_sequences.

Sequence 554 BP: 147 A; 100 C; 113 G; 154 T; 0 other;

Human genome-derived single exon nucleic acid probes "serul for analyzing gene expression in human placenta .

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Best Local Similarity 100.0i; P
Matches 25; Conservative 0;
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26 MAY 2000;
30 JUN 2000;
03 AUG 2000;
21 SEP 2000;
27 SEP 2000;
04 OCT 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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genetic disorder; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI39509 standard;
                                   The present invention relates to single exon nucleic acid probes (SEAr). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placents. The probes are useful for antenatal diagnosis of human genetic disorders.
                                                                                                                                                                                                                                                                                                                                          (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US00663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200157272-A2.
                                                                                                                                                            Claim 25; SEQ ID No 8195; 654pp; English.
                                                                                                                                                                                                                                                              WPI; 2001-488897/53.
                                                                                                                                                                                                                                                                                                      Penn SG,
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                                                                                                                                                                                                Human genome-derived single exon nucleic acid probes useful for
analyzing gene expression in human placenta .
                                                                                                                                                                                                                                                                                                 Hanzel DK, Chen W, Rank Dk;
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2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0236359.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; placenta; antenutal diagnosis;
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                             cc present sequence is human glycosyl sultotransferase-6 (GST-6)
cc percent DNA
cc percent suppe 2 meatrane protein useful for inhibiting a Linding e.e.f.
cc GST is a type 2 meatrane protein useful for inhibiting a Linding e.e.f.
cc between a selectin and a selectin light, which comprises contacting the
cc selectin with a non-sulphation activity of GST and a small molecular
cc agent that inhibits the sulphation activity of GST is also useful
cc agent that inhibits the sulphation activity of GST is also useful
cc in inhibiting a selectin mediated Linding event. GST is also useful
cc in inhibiting a selectin mediated Linding event. GST is also useful
cc in inhibiting a selectin mediated core chronic inflammation.
cc therapy to treat disorders such as acute of chronic inflammation.
cc incluse, polymyositis, dermatcupyositis, systemic salutoris, diabetes.
cc incluse, polymyositis, dermatcupyositis, systemic salutoris, diabetes.
cc glomerulonephritis, myasthenia gravis, Sjogren's syndrome, Hashimoto's
cc disease, Grave's disease, actematitis, hypocranthyroidism, permicious
cc anaemia, demyelinating diseases, cirrhosis, ulcerative collitis,
cc syndrome, infantile eczena, psoriasis lichen planus, aliergic rhinitis.
cc bronchial asthma, hypocrasnsitivity, rheumatic fever and tissue rejection
xx
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New glycosyl sulforrensferases (GSI)-talpha, GST-theta and GST-6 to diagnostic and therapeutic agent screening applications .
                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 116-123; 12spp; English.
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                             Sequence 27150 BP: 8357 A; 5396 C; 5398 G; 7998 T; 1 other;
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Query Match 2.5%; Sometimes Sest Local Similarity 100.0%; I Matches 25; Conservative 0;

Score 25; Pred. No. Mismatches

DB 22; Length 554; 0.72; ٥. Indels

0;

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RESULT 20 AAZ93815

AA293815 Standard; DNA; 144460 BP

AA253615;

Query Match 2.5%; Best Local Similarity 100.0%; Matches 25; Conservative (

Score 25; DB 22; ); Pred. No. 0.46; 0; Mismatches 0;

0; Indels 0; Length 27150;

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0:

AAD02701 standard; DNA; 27150 BP

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Human; glycosyl sultotransferase-6; GSI-6; immunosuppressive: therapy; selectin binding inhibitor; gene therapy; inflammation; systemic thuse brythematosus; SEE; rheumatoid arthritis; dlabetes; polyarteritis nodosa; polymyositis, systemic scherosis; dermatitis, glocarulonephritis; myastheria gravis; SJogren's syndrome; adrenaltis; Hashinoto's disease; chave's disease; hypoprathyroidism, anæmia; myocarditis; adult respiratory distress syndrome; accema; psortasis; sesthma; hypersensitivity; rheumatic fever; tissue rejection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human glycosyl sulforransferase-6 (GSI-6) genomic DNA #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62-MAY-2001 (first entry)
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                                                                                                                               20-JUL-1999; 9503-0144694
13-JUL-2000; 2000US-0593828
                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                               19-JUL-2000; 2000WO-US19741
                                                                                                                                                                                                                                                                                                 25-JAN-2001
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kosen SD, Lee JK, Hemmerich S;
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                                                                                                                                                                             /*tag- j
/label- ORF10
/label- ORF10
complement (113641..114063)
/*tag- k
/note- *Ubiquitin 1 pseudoyene complement*
complement (127469..127854)
/*tag- 1
/note- *Ubiquitin 2 pseudoyene complement*
7521
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110758.111564
110758.111564
110758.112267
122525.122667
12452.133369
12454.1133369
*tag 0RF9
123454.1133369
*tag 0RF9
113398.143577
                                                                                                                                                        /*tag- m
/note- *Polymorphic base G or T*
6192
                           /note- 'Polymorphic base A or C' $1088 /*tag- s
/note- 'Polymorphic base G or C' $1138
      /*tag* t
/note* "Polymorphic base A or G"
91187
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label- ORF 5
6291..96902
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711. 10658
**Lag - ORF |
10658
**Lag - ORF |
10658
**Lag - ORF |
10661 - ORF |
                                                   *Polymorphic base A or C*
                                                                       q
*Polymorphic base A or G*
                                                                                            p
*Polymorphic base C or T*
                                                                                                                 o
Polymorphic base A or I
                                                                                                                                      n
"Polymorphic base G or T"
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PT	PI	××	DR	××	Id	××	FA	XX	PX	XX	4.3	XX	Dd	××	PN	XX	F1	FI	1.3	FI	F.7	1.3	FI	FI	FI	FI	1.3	1.1	F) 7
and their biallelic markers, are useful in genetic analysis and in	New fucters actus encoding ten different olfactory receptor proteins		WPI: 2000-317933/27.		Bougueleret L,		(GEST ) GENSET.		14-CCT-1558;		13-OCT-1999;		20-APR-2000.		WU200021985-A2.				aliele			allele			allele			allele	
lelic mar	ius encod		33/27.		Malekzadeh K;		•		98US-0104299.		55WO-IB01729.				•		/note-	/*tag-	144033		/*Lag-		/1.0te-	/*tag-	140066				
kers, are us	ing ten diff				ideh K;				14259.		1729.						/note~ "Polymorphic base A or G"	Y		"Polymorphic base & or G"	×		*Polymorphic base C or T*	£		*Polymorphic base A or G	۷.		*Polymorphic base A or G*
eful i	tent																base			Lase			Lase			DáSe			LaSe .
g G	0116																۶ ۲			5			15.0			7.0			70.4
enetic	actory																6			G						6			G.
analysis	receptor																												
and in	proteins																												

Claim 1; Page 103-141; 155pp; English. screening for compounds which bind to the receptor proteins

Ten new offactory receptor proteins and their biallelic markers have teen described. The sequences encoding these receptor proteins and which contain the biallelic markers can be used for genotyping. The olfactory receptor proteins can be used to screen for substances which bind to them. See GENESED records AA253816:25 and AAY83386-95.

Sequence 144460 BP; 46068 A; 27088 C; 26615 G; 44676 T; 13 other;

£ Ŷ Ouery Match. 2.5%; Score 25; DB 21; Length 144460; Best Local Similarity 100.0%, Pred. No. 0.38; Matches 25; Conservative 0; Mismatches 0; Indels 0; 0;

RESULT 21 AAH51601 ID AAH516 XX Human chromosome 13q31-q33 genomic nucleotide sequence. 29-AUG-2001 (first entry) AAH51601 standard; DNA; 319608 BP.

sbg1: g34665; sbg2; g35017; g35018; chromosome 13g31-q33; haplotype; biallelic marker; polymorphism; schizophrenia; bipolar disorder; ds.

30-MAR-2000; 2000WO-IB00435.

05-001-2000

K0200058510-A2.

Gaps Ċ

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RESULT :
AASO301L ID
ID AASO301C AASO
XX AASI
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DT 26:
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                                                                                                                                                                                                                                                                                                                                                                              AAH51601 represents a human genomic nucleotide sequence comprising sbyi.

G 93466, sby2, 93017 and 935018 mucleic acid sequences located on the
Chuman chromosome 193119 locus. The nucleotide sequences contain
C biallelic markers and polymorphisms. Sequences AAH51602 AAH51620 AAH51631 represent cDNA human sbyl cDNA sequences and protein
C products. AAH51627 AAH51631 and AA562916 AAH51652 AH481652 are used
C to isolate sbyl cDNAs, while sbyl exons from different primates are
C tropresenced by sequences AAH51622 AH81669 Nucleotide sequences of
C amplicons which comprise biallelic markers located on the chromosome
C 1331-131 locus are represented in AH51700 AH81617. Hallelic marker
C are represented by adjunces by degenerate/undafined base codes. PCR
C tropresented in the sequences by degenerate/undafined base codes. PCR
C are represented in AH518181 are used in the isolation of sequences are
C used to determine the identity of the nucleotide at a biallelic marker in
C a sample DNA sequence. The nucleotide sequences may be labeled and used
C a sample DNA sequence. The nucleotide sequences may be labeled and used
C a sample DNA sequence. The nucleotide sequences are considered by determining the identity of the sucleotide at a biallelic marker in
C population an association between a genergy of a biallelic marker in a protein of the disorder or a beneficial response to or side cifects related to
C used to determine a predisoposition to or early onset of schizophrenia or
C used to determine or or a beneficial response to or side cifects related to
C treatment against schizophrenia or bipolar disorder.
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Best Local S
Matches 25
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30-APR-1999;
10-APR-1999;
11-JUL-1999;
27-JUL-1999;
29-JUL-1999;
29-JUL-1999;
29-JUL-1999;
                                                             Human; g35030; biallelic marker; al-A71; chromosome 13qil-q33; schizophrenia; bipolar disorder; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 409-493; 737pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polynucleotides comprising sequences from sbjg and 95508 bhallelic markers are used for genotyping and detecting schizophrenia or bipoiar disorder and predisposition to these disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cohen D. Blumenfeld M. Chumakov I. Bougueleret L. Essioux L:
                                                                                                                                Human schizophrenia associated gene g35030 and biallelic markers Al-A71.
                                                                                                                                                                                                                                                                      AAS09301 standard; DNA; 319608 SP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-615082/59
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al Similarity 100.0%;
25; Conservative 0
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990S-0131971.
990S-0132065
990S-0143928
990S-0146452
990S-0146453
990S-0146453
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/note* *Biallelic marker Al*
complement (8317..6335)
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/note= *Binds primer 59-27543-150.mis*
8304..8328
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/note- *Binds primer $9·31960-363.mis complement*
complement (95543..95563)
/*tag- x
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55377..95395
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/note* *Binds primer 6-128-33.mis complement*
complement (65856..65874)
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/note* *Biallelic marker A3*
complement (65486..65504)
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complement (8446..8465)
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7938..7958
                                                                                                          /note- "Biallelic marker A4"
complement (95397..95415)
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95034..95053
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21365..21385
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53..21671
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e+ *Biallelic marker A2*
lement (21673..21691)
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3..65471
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ement (21845..21864)
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...65497
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*Binds primer 8-128.pu*
..65484
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misc_binding
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                                            /"ay" au
/note= "Binds primer 99-7652-162.mis complement"
complement (169331..169351)
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ement (161240..161257)

** Binds primer 59.24634.rp complement*
3..168830
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= "Biallelic marker A5"
ement (107282...107300)

    "Biallelic marker A6"
    ement (168975...168993)

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4..160888
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ement (160785..160802)
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ement (160641..160659)
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*Binds primer 99-7652-162.mis*
..166986
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Binds primer 99-7652.pu*
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"Binds primer 99-24639.pu complement"
..160875
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*Binds primer 99-24639-163.mis complement*
...160787
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ment (107495..107513)
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..107280
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*Binds primer 99-7652.rp complement*
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                              21-FEB-2000; 2000EP-0200610.
                                                                                  06-SEP-2000
                                                                                                                                      EP1033401-A2
                                                                                                                                                                                                                                  Human; 5' EST: expressed sequence tag: secreted protein; cDNA isolation;
gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                              Human secreted protein 5' EST, SEQ ID NO: 32377
                                                                                                                                                                                                                                                                                                                                                                  06-CCT-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC26302 standard; cDNA; 305 BP
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//note* Biallelic marker Al0*
/note* Biallelic marker Al0*
complement (173359.173377)
/*tag* bg
/note* Binds primer 59-5862-167.mis complement*
complement (173495.173514)
/*tag* bh
/note* Binds primer 59-5862.pu complement*
189753..189771
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/*tag = a'

/*tag = a'

/*note* *Binds primer 59-16100.pu*

/*tag = a'

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18938. 18956
/*tag= bj
/note* "Binds primer 99-5919-215.mis"
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te* Binds primer $9-5862-167.mis*
346.173370
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*Binds primer 99-5862.rp*
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16

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Query Match 2.4%; So Best Local Similarity 100.0%; P Matches 24; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is one of a large number of 5: ESTs derived from mRNAS encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5 ESTS were prepared from the present sequence. The 3 ESTS were prepared from the present sequences usually correspond mainly to the 3 unitaristated region (UTR) of the mRNA because they are often obtained from obtained romalized region (UTR) libraries. Such ESTs are not well saited for isolating cDNA sequences they been obtained, the full 5 UTR is rarely included. 5' ESTS are derived from mRNAS and even in those cases where longer CDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTS are derived from mRNAS with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAS. 5' ESTS are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors.
                                                                                                                                                                                                                                                                                                                                                  Human; calcium sensitive potassium channel; beta2 subunit; asthma; beta3 subunit; diabetes; chromosome 193-ter; inhibitor; activator; glaucoma; migraine; angina; irritable bowel syndrome; Alzhelmer's disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID 32377; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid that is a 5' expressed sequence tag (5' ESI) for obtaining cDNNs and genomic DNNs that correspond to 5'ESIs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-MAY-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF27996 standard; DNA; 48000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dumas Milne Edwards J. Duclert A. Giordano J:
Novel human calcium sensitive potassium channel subunits for
                                                                           Uebele V, Swanson R, Liu Y, Lagrutta A;
                                                                                                                     (MERI ) MERCK & CO INC.
                                                                                                                                                                                                18-JUL-2000; 2000WO-US19585
                                                                                                                                                                                                                                       25-JAN-2001.
                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human calcium sensitive potassium channel beta3 subunits coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 305 BP; 80 A; 32 C; 68 G; 124 T; 1 other;
                                                                                                                                                                                                                                                                         WO200105828-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9905-0144764.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24: DB 21; Length 305; Pred. No. 1.9; O; Mismatches O; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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misc\_reature

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/label EST\_matching\_region
/label EST\_matching\_region
/note "This sequence is specifically claimed'
28364..28405

/note\* \*Interval associated with neuropsychiatric disorders\*
28441..29265

misc\_feature Homo sapiens

Location/Qualifiers 28441..144419

satellite primer\_bind

/rpt\_type= TANDEM /note= "BADCt22 marker" complement (28547..28572) /label\* BADct22\_forward 28547..28572 /\*tag= d

misc\_feature

7 del 9 /label EST\_matching\_region /note= "This sequence is specifically claimed" 43518..46075

/label- EST\_matching\_region /note- "This sequence is specifically claimed" 40284..43253

misc\_feature

misc\_feature primer\_bind

\* BADct2\_reverse .39587

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Query Match 2.4%; So Best Local Similarity 100.0%; I Matches 24; Conservative 0;
                                                                                                                                                                                                               The present invention provides the protein and coding sequences of the human calcium sensitive potassium channel beta?, beta3, beta3 beta3b, beta3f activators in these can be used to identify inhibitors and activators of the channels, which can be used in the treatment of conditions including astima, disbetes glucoma, cerebral ischaenia Alzheimer's disease, excessive smooth muscle tone, angina, hypertember incontinence, migraine and irritable bowel syndrome. The coding sequences are found at ruman chromosome 323 tter. The present sequence is the beta3 subunits coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; Fig 6; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       learling conditions such as astima, hypertension, memory disorders, depression
                                                                                                                                                                               Sequence 46000 BP; 13303 A; 9551 C; 10282 G; 14832 T; 32 other;
                                                                             Score 24; DB 22;
Pred. No. 1.1;
0; Mismatches 0
                                                                                  0; Indels 0;
                                                                                                                     Length 48000;
                                                                          Gapo
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RESULT 25 AASÖ4858/c ID AASÖ4858 standard; DNA; 160271 BP. himan; chrisq; rsh10; bipolar affective disorder; BAD; neuropsychiatric disorder; antibody; schlzophrenia, Alzheimer's dis. 13e; Hintitgdon's disease; Parkinson's disease; amyotrophic lateral sclessasis; 07-SEP-2601 (first entry) Human chromosome 18, BAD18ct22-BAD18cagl

	1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1	2.1. 2.1. 2.1. 2.1. 2.1. 2.1. 2.1. 2.1.		4 4 4 4 4 4	# # # # # # # # # # # # # # # # # # #		11111	11111	111111	7777	1333
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/*tag* X /label EST_matching_region /note* "this sequence is specifically claimed* 80466.81515 /*tag* Y /label* EST_matching_region /note* "This sequence is specifically claimed* 81888.85346	/*tag- v //tag- v //tag- v //tabel- ESI_matching_region //tag- v //tabel- ESI_matching_region //tag- v //tabel- ESI_matching_region //tag- v //tabel- ESI_matching_region //tag- v //tabel- v //tag- v //tabel- v //tag- v	/*tag- t. //tabel- EST_matching_region //abbl- EST_matching_region //note- *This sequence is specifically claimed* //tag- u. //tag- u. //tabel- EST_matching_region //note- *This sequence is specifically claimed* //199777123	/noce. "his sequence is specifically claimed" 70643.70749 /*tag	sequence is specifically	/*Lay- (1-14) / *Lay-	/label- EST_matching_region /note- "This sequence is specifically claimed" 59815. 6071 /tag- n /label- EST_matching_region /note- "This sequence is specifically claimed" 60870 62487	/tag. 1 /tag.	520725935  /*tay* k /label* EST_matching_region /note* "This sequence is specifically claimed* 57010 57705	47645284  /*tag	/*tay . 1 - 40//5 /*tay . 1 /*tay .	/*tag. h /label. ESI_matching_region /note. This sequence is specifically claimed.
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RESULT 26

ANSO4864;

TD AASO4864;

XX AASO4864;

XX AASO4864;

XX CAASO4864;

XX CAASO4864;

XX AASO4864;

XX Human chromosome 18, BAD18ct22-BA
DE Human chromosome 18, BAD18ct22-BA
XX Human; chr18q; fsh27; bipolar aff
XX Human; chr1
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neuropsychiatric disorder; antibody; schizophrenia. Alzhenter's disease;
Huntinydon's disease; Parkinson's disease; amyotrophic lateral sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human chromosome 18, BAD18ct22-BAD18cag1.
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131138...134228
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28441.144419
/*Lag* a
/mote* "Interval associated with neuropsychistric
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43518..46075
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complement (28547..26572)
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note- "This sequence is specifically claimed"

10284...43253
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note- "This sequence is specifically claimed"
8384..28405
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3518..46075
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3547..28572
"This sequence is specifically claimed"
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52672...5935
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/note* This sequence is specifically claimed*
86346.87569
/1eg* aa
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/label- ESI_matching_region
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78463. 80173
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| 11888..85946
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/note* "This sequence is specifically claimed"
77663..78170
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70643..70749
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89459..85745
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101798..103865
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101798. 103865
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note* "This sequence is specifically claimed'
.00530...101382
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116846...117907
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'label = EST_matching_region

note "This sequence is specifically claimed"

0953 ...10561
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13774..116253
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11000..113482
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17999. 118623
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5814..119863
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Matches 24
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misc\_feature 1.cm 2.4%; Score 24; LB 22; Length 160271; al Similarity 100.0%; Pred. No. 0.94; 24; Conservative 0; Mismatches 0; Indels 0; 131136...34226
/\*Lay- at
/label- EST\_matching\_region
/note- \*This sequence is specifically claimed\* Gaps

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Human, 18q; fsh.3; neuropsychiatric disorder; schizophrenia; neurobagenerative disease, Alzheimer's disease; Parkinson's disease; Erain tumour; diseases; angina paecoris; ds. Human chromosome 18q, 160kb sequence. AAS06667 standard: DNA; 160271 SP 12-SEP-2001 (first entry)

misc\_feature misc\_feature Homo sapiens. primer\_bind /note- "This region is specifically claimed" 25683.35587 /note= "PCR primer BAD18ct22 forward" 28441 29265 /\*tag- b Location/Qualifiers 28384..28405 "This region is specifically claimed"...144415

misc\_reature misc\_reature primer\_bind misc\_feature /\*tag. fi note. This region is specifically claimed. 1518. 16075 /\*tag. 7his region is specifically claimed. 17264. 52264 /\*tag. fi /\*tag. fi /\*tag. i /\*Lag\* e /note\* \*PCR primer BAD18ct22 reverse\* 40284..43253 complement (28547..28572) note= \*Region associated with neuropsychiatric

misc\_teature misc\_reature misc\_feature misc\_feature /note- This region is 59815..60471 /note- \*This region is 58065..59057 /note- "This region is specifically claimed" 57032..57726 /product\* "Fsh23" /note\* "Fsh23 gene region" 52672..56935 region is specifically claimed\* specifically claimed. specifically claimed.

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                                          C. file Sequence represents 160kb of human chromosome 18g containing C. file3 yene, locked in a region associated with neuropsychibution of obscrates. The file2 yene and polypeptide, its iragment, analog or callectic is useful for treating a file2-related disorder or file3-mediated electric is useful for treating a file2-related disorder. Dipolar Affective attention derical disorder, schizopireder, bipolar Affective disorder and unipolar disorder. A cell harbouring the gene and protein are calso useful for treating neurodeptimentive gene and protein are calso useful for treating neurodeptimentive disorder such as Atzehner's calso useful for treating neurodeptimentive disorder such as Atzehner's calso useful for treating neurodeptimentive disorder such as Atzehner's collects, parkingon's disease, and cilies de la Tourette's syndrome, and typertension and sleep disorders, mania, cusessive compulsive disporter. C. Expertension and sleep disorders, mania, cusessive compulsive disporter. C. evaluation, genetic testing and/or prognosis of a fsh23-related disorder, evaluation, genetic testing and/or prognosis of a fsh23-related disorder.
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(REGC ) UNIV CALIFORNIA.
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Sequence is only 160kb in size.
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This invention relates to the present sequence which represents a look fragment of the long arm of human chromosome ldq. The sequence includes the 18q interval associated with neuropsychiatric disorders (unclavidus 28441-1444)9). Included in this region is the fsh2 gene, which is involved in neuropsychiatric disorders. Included in the invention arc
                                                                                                                                                         Claim 1; Fig 1B; 153pp; English.
                                                                                                                                                                                                         Nucleic acids encoding mammalian rsh25 polypeptides ussociated with neuropsychiatric disorders, useful for treating e.g. bipolar affective disorders.
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/product= "fsh25 protein"
complement (144302..144325)
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complement (28384..28405)
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complement (116254..116523)
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28547..28572
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28440..28483
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"BAD18cag1 specific primer"
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   RESULT 29
AH23764/C
AH23764 standard: DNA: 160271 BP.
ID AH23764:

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XX AH23764:

XX Human chromosome 18q: fsh26; nootro
KH neuroleptic: hypotensive; gene thera
KH neurolepsychiatric disorder; senie thera
KH neurolepsychiatric disorder; senie thera
KH neurolepsychiatric disorder; senie thera
KH anyotrophic lateral sclerosis; Parki
KH Gilles de la Tourette's syndrome; ds
XX Location/Qualifiers
FI misc_feature 25683.3567

FI misc_feature 47264.3253

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FI misc_feature 57032.5294,

FI misc_feature 57032.5726.

FI misc_feature 57032.5726.

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Best Local Similarity 100.0%;
Matches 24; Conservative (
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25683..39587
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28441..29265
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/note= This sequence is
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/note= "This sequence
113774..116253,
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/note- "This sequence
111000..113482,
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100530..101382,
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note- "This sequence
04486..109841,
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note= "This sequence
09953..110561,
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131136..134226,
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/note= *This sequence is
129508...130413,
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116665..122881.
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This sequence is specifically claimed in Claim 1.
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...135473,
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17 - MAY - 2001

ŭ8-NJV-≥₀∪0; 2666NJ-8530824

08-NOV-1959; 990S-0164037.

(MILL-) MILLENNIUM PHARM INC (REGC ) UNIV CALIFORNIA.

Chen H, Freimer NB

WPI; 2001-335946/35.

I mamalian (sh/6 polynoclevide for diagnostic evaluation, genetic testing and prognosis of isb/6 related discretes e.g., neuropsychiatric discretes including schizophrenia and hipolar affective disorder

Claim i; Fig 1B; 174pp; English.

The present sequence is a 116 kb fragment located between markers CR abbiect22 and Bablecapi from human chicaccame 184, This sequence includes the 164 interval associated with neuropsychiatric disorders, located from positions 26441-144419. This sequence also contains a novel gene: 184.20 capae. The sinh5 gene is associated with bipolar affective disorder (BAD) and the present sequence or its fragment analog or minetic is useful for treating a fish5-related disorder: 54; or manic-depressive illness) in contains. The present sequence or its fragment analog or minetic is useful for treating a fish5-related disorder or ish5-mediated process in a cannal, such as neuropsychiatric disorders with as schloophrenia.

CR alterial such as neuropsychiatric disorders such as schloophrenia.

CR disorder and unipolar disorder. Also, neuropsychiatria disorder bipolar affective disorder bipolar affective disorder bipolar affective disorder bipolar affective disorder. Such as a schlophic lateral scherosis, and Parkinson's disease, as vell as Gilles de la curette's syndrome, autonomic function disorders such as hypertension cand sleep disorders can be treated.

Sequence 160271 BP: 45618 A; 32564 C; 34528 G; 46703 T; 58 other:

Query Match 2.4%; So Best Local Similarity 100.0%; 1 Matches 24; Conservative 0; Score 24: LB 22; Pred. No. 0.54; 0; Mismatches 0 0; Length 160271; Indels 0 Gaps

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misc\_feature

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AAF83116/C

AAF83116:

XX

AAF85116:

XX

Muclectide sequence of 160 kb of chr

XX

BAD18cag1 marker; fsh28 gene; bipola

KW chromosome 18; chromosome 18q; neuco

KW unipolar disorder; BAD18ct22 marker;

XX

HOMO sapiens.

XX

AND 17-AAY-2001.

XX

AND 17-AAY-2001.

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AND 18-AAY-2001.

AND 18-AAY-200
                                                                                                                                                                                                                                                   RESULT 31
AAF93676
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Best Local Similarity 100.0%;
Matches 24; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents a 150 kb fragment of the long arm of chromosome 18. This region is associated with neuropsychiatric disorders, and contains the fishal gene and markers subject2 and BAD18649. The fishal gene is a gene involved in bipolar affective disorder (BAD). The fishal polyniciatides and polypeptides are used to detect, modulate, or reast chromosome 18g related disorders are used to detect, and the subject of the first polynemia, attention deficient disorder; a schizopared attention deficient disorder, a schizopared can be used to identify an individual having or at risk of developing one of the disorders. The fishal gene and derived polynicipations can be used to identify a compound capable of modulating fishal activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BAD18cagl marker; fsh28 gene; bipolar affective disorder; BAD; chromosome 18; chromosome 18q; neuropsychiatric disorder; schizophrenia; attention Jeficient disorder; schizoaffective disorder; unipolar disorder; BAD18ct22 marker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated mammalian nucleic acid for disposing, monitoring and treating neuropsychiatric disorders, such as bipolar affector disorder, schizophrenia, or attention deficient disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleotide sequence of 160 kb of chromosome 18q
                     cDNA isolated from foetal brain tissue encoding SRF protein SEQ ID 497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 160271 BP; 45619 A; 32964 C; 34928 G; 46702 T; 58 other;
                                                                                               21-MAY-2001 (first entry)
                                                                                                                                                                                                                         AAF93676 standard;
                                                                                                                                                                                                                         CDNA; 553 BP
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KESULI 32

AAC578917C

ID AAC57891 standard: LNA: 1001 EP.

XX AAC57891:

AC AAC57891:

XX AC57891:

XX Arachidonic acid metabolism relate

XX Arachidonic acid metabolism relate

XX Metabolic marker: arachidon

XX Human: biallelic marker: arachidon

XX detection; hybridisation; phenotyp

XX single nuclection of assay; iden

XX specific amplification assay; iden

XX Homo sapiens.

XX Mc200047771-AZ.

XX Mc200047771-AZ.

XX D1 AUG-2000.
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Query Match 2.3%; Best Local Similarity 100.0%; Matches 23; Conservative (

0;

Score 23: LB 22: Length 553; Pred. No. 4.4; 0: Indels

0 Gaps

0

Sequence 553 EP; 156 A; 108 C; 51 G; 198 I; 0 other;

Human; biallelic marker; arachidonic acid metabolism; genotyping; detection; hypridisation; phenotype; haplotype; SNP; polymorphic base; single nuclectide polymorphism; hybridisation assay; sequencing assay; specific amplification assay; identification; ERBH; 12:10-RBH; eccessicid-related biallelic marker; 12:10-related biallelic marker; ds.

Arachidonic acid metabolism related genomic biallelic marker #525

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Sequences AAF93180 - AAF93741 represent polynucleotide sequences encoding human SRI proteins. The cDNA sequences are isolated from various CC different human tissue cDNA libraries. The invention relates to a method of iferent human tissue cDNA libraries. The invention relates to a method conding SRI, a host cell transformed with the vector containing cDNA. CC encoding SRI, a host cell transformed with the vector, an isolated SRI colyopepide, and an anthody which binds to SRI. The polynucleotide sequence can be used in gene therapy and is useful in the recombinant production of SRI polypepides, as a hybridisation probe to screen composition of SRI polypepides, as a hybridisation probe to Sereen colors to isolate cDNAs with sequence identity to SRI polypepides, to a libraries to isolate cDNAs with sequence itssue detection. The SRI collisorders, tissue typing and disease tissue detection. The SRI collisorders, tissue typing and disease tissue detection. The SRI collisorders, tissue typing and disease tissue detection. The SRI collisorders, tissue typing and disease tissue detection. The SRI collisorders, tissue typing and disease this reaction.
                                                                                                                                                                                                                                                                                                                                                                                                            New isolated nucleic acid molecule encoding a SRT polypeptide is for production of recombinant SRT polypeptides, gene mapping, diagnosing genetic disorders and for gene therapy
                                                                                                                                                                                                                                                                                                                                                                     Claim 2: Fig 457; 663pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    baker KP, Goddard A, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GEIH ) GENEMIECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-112729/12.
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12-FEB-1999; 23-MAR-1999; 07-MAY-1999;

9905-0115917. 9905-0275267. 9505-0133200.

(GEST ) GENSET.

11-FEB-2000; 2000WO-IB00184

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RESULT
AAD04475
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OR M.B. Polymarphic bases (single nucleotide polymarphisms also known as the polymorphic bases (single nucleotide polymarphisms also known as constituted programment invention have been given as their corresponding degenerate bases e.g. a polymorphic base of C or T has been given as X.
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Best Local Similarity 100.04; E
Matches 23; Conservative 0;
                                                                                                                             Human; ADAM-TS; A Disintegrin And Metalloproteinase; antiinflammatory; anglogenesis; bronchial astima; Goodpasture is syndrome; metastesis; heart failure; cardiac hypertrophy; chronic ischaemic heart disease; sickle cell disease naphropathy; urinary tract obstruction; haemostatic; skeletal muscle disorder; myocardial infarction; blood vessel disorder; myocardial infarction; blood vessel disorder; myocardial infarction; blood vessel disorder; sonorrhea; tuberculosis; vasculitis; renal artery santoma: osteoporosis; mumps; gonorrhea; tuberculosis; syphilis; spermatocytic santoma: osteoporosis; ribetomyosarcoma; glomerulonephritis; bone disorder; baget's disease; ribtets; osteomalacia; hodsylin; od disease; gono therapy; antibacterial; cardiant; tumour; thymoma; vasouropic; cytostatic; virucide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 13; Page 673; 802pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel biallelic markers useful for detecting conditions and genotypes associated with arachidonic acid metabolism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-571881/53.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      04-JUL-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD04475 standard; cDNA; 5353 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1001 BP; 305 A; 182 C; 279 G; 254 i; 1 other;
                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                        Human 27875 ADAM-TS cDNA, alternative version.
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Location/Qualifiers
36..5096
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Pred. No. 4.1:
0: Mismatches 0: Indels (
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Kovel isolated golypeptide, 27875, a human ADAM-TS (a disintegrin and becalloproteinuse) useful for diagnosis and treatment of disorders of boes, lung, heart, skeletal muscle such as osteoporosis, emphysema, anglica.
                                                                                                                                                                                                                                                                                                                                                                                                 sig_peptide
                                                                                                                                                                                                                                                                                                  (MILL-) MILLEGNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                              25-CCI-2000; 2000WO-US29360
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P-PSDB; AAE00913.
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                                                                                                                                                                                                                                                                                                                                                          ⊮∪200131v34-A1
                                                                                                                                                                                                                                                                                                                                                                      /product. *Human 27875 ALAM-TS protein*
/transl_except. (pos:1e21..182), aa:Xaa)
/cote. *Xaa is an unknown amino acid*
56..155
/*tag. b
/*tag. b
/*tag. c
/product. *Human mature 27875 ALAM-TS protein*
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Claim 7; Fage (15-123; 133pp; English.

The present separation and alternative version of a cDNA encoding C2 7875 protein, a miman ADAM-TS (a disintegrin and metalloproteinase). The capturation, protein degradation, tumbur growth, metalsais and C2 capturation, protein degradation, tumbur growth, metalsais and C3 capturation, protein degradation, tumbur growth, metalsais and its C3 antitudates are usually for preventing, dispinosing and treating C2 antitudates are usually for preventing, dispinosing and treating C3 include, disorders involving the luma such as congenital anomalies. C3 include, disorders involving the luma such as congenital anomalies. C4 disorders involving the skeletal muscle include tumours such as bronchial astlam, coopsasture's synorame, pulmonary alveolar proteinosis. C4 disorders involving the skeletal muscle include tumours such as capture such as theroscies involving the skeletal muscle include tumours such as capture and storders involving blood vessel such as hypertension, captured the state and geldidynis such as onespecific epidolymitis and crecities, paper structure, disorders and syphilis, spermatocytic seminours put the testis and geldidynis such as onespecific epidolymitis and crecities, paper sinvolving the kidney such as membrangoroliferative c3 captacosis, circuit glomerulonephrittis, sicke cell disease rephropatry, carlocals, circuit glomerulonephrittis, sicke cell disease rephropatry, carlocals, capture solution, osteonar come, metastatic tumours, call disorders involving thymus such as athomatory tumours, and disorders involving thymus such as trymic bypoplasia. C5 captures shown in Fig 1 (See ADO4494), However these sequences ditter at several locations.

Sequence 5353 6P; 950 A; 1833 C; 1648 G; 911 T; 1 other;

Guery Match 2.3%; Score 23; DB Best Local Similarity 100.0%; Pred. No. 3.4 Matches 23; Conservative 0; Mismatches DB 22: Length 5353: 0; Indets 0 Gaps

į,

04-JUL-2001 (first entry)

Human 27875 ADAM-TS (a disintegri. and metalloproteinase) cDWA

AAD01191 standard;

CDNA; 5353

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The present sequence is a cDNA emining 27875 protein, a human ADAN-IS CC (a disintegrin and metalloprotein:se). Metalloproteinse is a proteolyptic earyme involved in purchin maturation, protein degradation, cc tumour growth, metastasis and analysagenesis. Nucleotides encoding 27875.

CC 27875 protein and its antibodies are useful for preventing independency include, disorders involving the include suseful for preventing independency considers involving the include suseful for preventing alveolar proteinset for an artificial stifma. Goodpasture's a mature trailure, cardiac hypertrophy.

CC disorders involving the stellar involving the preventing at recolar proteins. The disorders involving the stellar involving the stellar involving the stellar involving the trailure associated with other disorders disorders involving the testis and epididy, as such as nonspecific epididymitis and cribitis, somerine, mumps, tuther disors as sphills, spermacocytic glomerulophritis, percetsing involving such as another alsorders involving the stellar involving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ADAM-TS; A Disintegrin And Mctalloproteinase; antiinflammatory; andylogenesis; bronchial asthma: "... Appasture's Syndrome; metastasis; heart failure; cardiac hypertroph; chronic isohaemic heart disease; sickle cell disease nephropathy; ... innary tract obstruction; haemostatic; skeletal muscle disorder; myocariat infarction; blood vessel disorder; skeletal muscle disorder; myocariat infarction; blood vessel disorder; synphitis; renal artery stencesis; ... maps; gonorrhea; tuberoulosis; v.a. vilitis; renal artery stencesis; ... maps; gonorrhea; tuberoulosis; syphilis, spermatocytic senioma: osteoprosis; Tabadomyosarcoma; glomerulonephritis; bone disorder; Paget's disease; rickets; osteomalacia; Hodykin's sisease; gane threopy; anticacterial; cardiant; tumour; thymoma; vasot: ..., ic; cytostatic; virucide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated polypeptide, 2767: a human ADAM-TS (a disintegrin and metalloproteinase) useful for dispusis and treatment of disorders of bone, lung, heart, skeletal muscic such as osteoporosis, emphysema, angina
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 7; Fig 1; 133pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -PSDB;
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DB; AAE00934.
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36-KCV-2000. K3200071703-A2 nomo sapiens

J3-MAY-2000; 2000WO-IB01252

9905-0132287

Macleod AR, Li Z, besterman (METH-) METHYLGENE INC J3-MAY-1999;

JH;

2001-016407/02. UB; AAB49559.

17 S

628 AALCIGIAGANIGCIIIGGGIAA 44091

Guery Match Best Local S Matches 23

ch 2.3%; Score 23; LB 1 Similarity 100.0%; Fred. No. 2. 23; Conservative 0; Mismatches

LB 22; 9

Length 60331; Indels

0:

Gaps

0

The present invention provides inhibitors of histone descriplase enzy, such as HDAC-1, HDAC-2, HDAC-3, HDAC-3, HDAC-4, HLAC-5, HDAC-C and HDAC-D This inhibitors may be antisense strands or they may be compounds identification of the enzyme with the compound and measuring the resulting concerts and identify, these inhibitors are useful for treating cancers and identifying which histone deacetylase is involved in a neoplasia.

:5: - 00 00

Sequence 60331 BP; 26046 A; 14486 C; 13823 G; 25976 T; 0 other;

Artiseise oligonopiocitàe that inhibits expression of a histone deacetylase, useful for treating acd/or alleviating the symptome acceptasia, or for inhibiting neoplastic cell growth in an animal

toms imal

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Disclosure; Page 66-89; 125pp; English.

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                                                                                                                                                                                             323333333
                                                                                 Every Match 2.3%; Sometive 100.0%; Fact Local Similarity 100.0%; Factories 23; Conservative 0;
                                                                                                                                                                                                                             Liberculous esteemyelitism, esteema, esteema metastatic tumouts and disorders involving thymus such as thymic cysts, thymic hympolasi, thymicans, and doogkin's disease. 27675 Sequence is used in gene there; // NCTE: This sequence SEG. ID. NO.2 is stated as being similar to the sequence SEG. ID. NO.2 is stated as being similar to see sequence slown in Page 115-123 (See AADG4175). However these sequences of the sequence slown in Page 115-123 (See AADG4175).
Sequence 5353 BP: 960 A; 1833 C; 1645 G; 911 T; 0 other;
                                                                                 Score 23; DB 22; Length 5353;
Pred. No. 3.4;
O; Mismatches O; Indels
                                                                                          0
                                                                                          Gaps
                                                                                             0
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RESULT 35 AA169559/c ID AAC89559 standard; DNA;

80331

g.

AAC69559;

us-MAR-2001 (first entry)

itstone deacetylase; HDAC-1; HDAC-2; HDAC-3; HDAC-4; HDAC-5; HDAC-C: HDAC-D: cell cycle; tumourigenesis; cancer; inhibitor; antisense; jene therapy; ds.

muman histone deacetylase HDAC-C coding sequence.

RESULT 36 AAC15149 ID AAC151

AAC15149 standard; cDNA; 247

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RESULT 37
AAC26972
ID AAC269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is one of a large number of 5 ESIS derived from CC mRRAS encoding secretch proteins. No ORF has yet been conclusively conclusively in the present sequence. The 5 ESIS were prepared from CC total human RRAS or polyar RNAS derived from 30 different tissues. ESI CC sequences usually correspond mainly to the 3 untranslated region (UTR) CC of the mRNA because they are often obtained from ollyor off primad cDNA CC libraries. Such ESIS are not well suited for isolating cDNA sequences (CC derived from the 5 ends of mRNAs and even in those cases where longer CC derived entry because they are often obtained. The full 5 UTR is rarely included. CS 5 ESTS are dot DNAs and appeare 5 ends of mRNAs with intext 5 ends and can therefore be CC used to obtain full length cDNAs and spannic DNAs. 5 ESTS are also used CC used to obtain full length cDNAs in a spannic by DNAs. 5 ESTS are also used CC in diagnostic, forensic, gene therapy and biromosome mapping procedures. CC They are used to obtain out to obtain unit of the procedures and contains and secretion vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 2.2%; So Best Local Similarity 100.0%; P Matches 22; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID 19224; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid that is a 5, expressed sequence tag (5, EST) for obtaining cDNAs and yenomic DNAs that correspond to 5 ESTs and for diagnostic, forensic, yene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-SEP-2000.
Human secreted protein 5' EST, SEQ ID NO: 31047.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dumas Milne Edwards J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-1599;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP1033401-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted protein 5' EST, SEQ ID NO: 19224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-OCT-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 247 BP; 55 A; 39 C; 51 G; 97 T; 5 other:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-500381/45.
                                                              06-OCT-2000 (first entry)
                                                                                                                            AAC26972;
                                                                                                                                                                                      AAC26972 standard; cDNA; 329 BP
                                                                                                                                                                                                                                                                                                                                   9905-0122487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Duclert A. Giordano J:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 22; DB 21;
; Pred. No. 12;
0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels 0:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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HLMan; 5' ES1, expressed sequence Ldg: Secreted protein; cDNA isolation: gene therapy; chromosome mapping: 55.
                                                                                                                                                                                                                     Dumas Milne nawards J. Duclert A.
                                                                                                                                                                                                                                                                                            06-SEP-20UU.
                                                                                                                                                                                                                                                                                                                               Homo sapiers
                                                                                                                                                   Claim 1; SEQ 10 31047; 71pp + CD-KGM; English.
                                                                                                                                                                  Kww fucleic acid that is a 5' expressed sequence tag (5' ESI) for containing cubas and genomic pakes that correspond to 5'ESIs and for diagnostic. forensic, space therapy and chromosome mapping procedures
                                                                                                                                                                                                      WP1; 2000-50usel/45.
                                                                                                                                                                                                                                       (GEST ) GENSEL.
                                                                                                                                                                                                                                                          26-FEB-1999;
                                                                                                                                                                                                                                                                          21-FEB-2000; 2000EF-0200610
                                                                                                                                                                                                                                                                                                             EP1033401-A2
                                                                                                                                                                                                                                                          59US-0122487
                                                                                                                                                                                                                         Giordano J;
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CC mkNAs encoling secreted proteins. No Dark has yet been conclusively complete the conclusively concentration of the concentration of

Sequence 329 BP; 80 A; 50 C; 66 G; 132 T; 1 other;

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Query Match 2.2%; So Best Local Similarity 100.0%; P Matches 22; Conservative 0:
Score 21, DB 21: Length 329;
Pred. No. 12;
0: Mismatches 0: Indels
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    Gaps
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9
                                                                                       93
                                                    01-FEB-2000
                                                           AAZ42729;
                                                                                      Human 5' EST isolated from a cDNA library SEQ ID NO:488.
                                                                   AAZ42729 standurd; cDNA; 329 BP
                                                                           36
                                                    (first entry)
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Human; 5: EST; expressed sequence tag; secreted protein; diagnosis; gene therapy; chromosome mapping; upstream regulatory sequence; forensis; location; development; protein synthesis; stability; regulation; luentification; ss.

21-001-1999

**W**09553051-A2

Homo sapiens

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A AAA42265 to AAZ43075 represent novel 5' expressed sequences tay (EST) sequences, corresponding to human secreted proteins. AAX6451 to AAX43052 the AAX64318 represent the EST-related proteins corresponding to AAX42265 to CAAX643052. The 5' ESTs can be used for producing secreted ...man gane corresponding to the AAX43052 they can be used to identify and isolate 5' untranslated regions (UTRs) and upstream regulatory regions which control the correct development stags, rate, and quantity of protein synthesis, as the correct development stags, rate, and quantity of protein synthesis, as covell as stability of mRMA. The ESTs are also useful as proteins for control comes mapping, and to obtain full length colk clones. The ESTs can disposite procedures to identify individuals having genetic diseases of the common sequences and the dentify individuals having genetic diseases cresulting from abnormal gene expression. The products may also be used in common sequences are perfectly of the colds encoding signal peptides on the close tracting a variety of human conditions. Secreted proteins the control of the proteins according to the conservation of a polypeptide of the insertion of a polypeptide into a membrane, or importing a polypeptide content of the proteins according to the conservation of a polypeptide of the proteins according to the conservation of a polypeptide conservation of a polypeptide of the proteins according to the conservation of the proteins according to the content of the proteins according to the conten
                                                                                                                                                                                                                                                                                                                                                                                                                            Owery Match 2.2%; Score 22; DB 21; Length 31; Best Local Similarity 100.0%; Prod. No. 12; Matches 22; Conservative 0; Mismatches 0; Indels
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28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 427; 837pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel secreted protein 5' expressed sequence tay sequences used in diagnostic, forensic, gene therapy, and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-038446/03.
P-PSDB; AAY65115.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dumas Milne Edwards J,
Arabidopsis thaliana DNA fragment SEQ ID NO: 13931.
                                                      17-OCT-2000 (first entry)
                                                                                                                    AAC36471;
                                                                                                                                                                           AAC36471 standard; DNA; 731 BP
                                                                                                                                                                                                                                                                                                               (GEST ) GENSET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Giordano
                                                                                                                                                                                                                                                                                                                                                                                                                                   ي.
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25-FEB-2000; 2000EP-0301439

06-SEP-2000. EP1033405-A2. Arabidopsis thaliana. Hybridisation assay, genetic mapping, gene expression varitul; protein identification, signal transduction pathway; metabolic pathway; promoter; termination sequence; ss.

Page

26

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PR 13-JUL 1999 99US 011542
PR 16-JUL 1999 99US 0114085
PR 16-JUL 1999 99US 0144085
PR 19-JUL 1999 99US 0144085
PR 19-JUL 1999 99US 0144313
PR 20-JUL 1999 99US 0144313
PR 20-JUL 1999 99US 0144314
PR 21-JUL 1999 99US 0144314
PR 21-JUL 1999 99US 014508
PR 22-JUL 1999 99US 014508
PR 22-JUL 1999 99US 014508
PR 22-JUL 1999 99US 014518
PR 22-JUL 1999 99US 014518
PR 22-JUL 1999 99US 014518
PR 22-JUL 1999 99US 0145218
PR 22-JUL
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Query Match 2.2%; Score 22; DB 21; Length 731;
Best Local Similarity 100.0%; Pred. No. 11;
Hatches 22; Conservative 0; Hismatches 0; Indels 0;

Gy 410 atalatatacacacatatatac 431

DD 393 atalatacacacacatataca 414
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Gaps

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AZZZEGZ

ID AAZZZEGZ

XX

XX

AAZZZEGZ

XZ

AC AAZZZEGZ:

LI 20-DEC-1999 (first entry)

XX

DE Kheat indole-3-glycerol phosphate synthase partial CDNA.

XX

KH Tryptophan; biosynthesis; transgenic plant; herbicide;

KH Indole-3-glycerol phosphate synthase; os.
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×		
S	Triticum aestivum.	YUM.
×		
H	Key	Location/Qualifiers
F	CDS	1.810
F		/*tag~ a
F		/partial
7		/product= "Wheat indole-i-glycerol phosphate synthase"
7		/note= "No initiation codon given in specification"
×		
z	W09949013-A2.	

26-MAR-1998;

9805-0079386

30-SEP-1999; 22-MAR-1999;

99MO-US06582.

(DUPO ) DU PONT DE NEMOURS & CO E I.

us-09-248-178-64.rng

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This sequence represents a wheat indole-3-glycerol phosphate synthase controlled to the biosynthesis of tryptophan. Plants are able to synthesise convolved in the biosynthesis of tryptophan. Plants are able to synthesise carlain anno acids, unlike vertebrate animals, which require controlled the manno acids (essential amino acids) in their diet as they cannot mannifecture them de novo. Tryptophan is one such essential amino acids (essential amino acids) in their diet as they caid, in plants, the biosynthesis of tryptophan is one such essential amino acid. In plants, the biosynthesis of tryptophan from chorismic acid requires five enzymatic steps catalysed by anthranilate synthase. Considerate, including the phosphate synthase and tryptophan synthase. Considerate, including the hormone insole-3-acetic acid, antimicrobial tryptophan biosynthetic enzymas may be used to generate transpanic content or those which cannot synthesise tryptophan biosynthetic enzymas is altered, including plants having an increased tryptophan content or those which cannot synthesise tryptophan (loss of function mutants). Content or those which cannot synthesise tryptophan (loss of function mutants). Content or specific inhibitors which may be potentially useful as herbicides. As these enzymes are involved in a biosynthesis path-sy that does not exist content or inhibitors which may be potentially useful as herbicides. As these enzymes are involved in a biosynthesis enzymes are succession or exist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cahoon RE, I
Rafalski JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 7; Page 55-56; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-580437/49.
P-PSDB; AAY42527.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated nucleic acids encoding plant enzymes involved in
tryptophan biosynthesis, used to generate transgenic plants, and
enzymes used as herbicide targets.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Falco SC, Gutteridge S, A, Tao Y, Vollmer SJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hitz WD. Maxwell CA:
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RESULT 41
AAC32953
TA AAC329
AC AC329
AC AC3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hybridisation assay, genetic mapping, gene express, protein identification, signal transduction pathway; mutabolic pathway; promoter; termination sequence; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arabidopsis thaliana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Arabidopsis thaliana DNA fragment SEQ ID NO: 1253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC32953;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels 0;
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25-FEB-1999; 05-MAR-1999;

99US-0121825 99US-0123180

25-FEB-2000; 2000EP-0301439.

06-SEP-2000. EP1033405-A2 В 2

Sequence 1005 BP; 282 A; 189 C; 270 G; 264 1; 0 other;

25 MAR 1955
25 MAR 1555
26 MAR 1555
27 MAR 1555
28 MAR 1555
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RESULT 42
AMH17506/C
AMH17506 standard: cDNA; 1796 BP
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AC AAH17506;
XC AAH17506;
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AC AAH17506;
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AC AAH17506;
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AC AAH17506;
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BE Human cDNA sequence SEQ ID NO:16
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BH.man; primer: detection; diagno
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BEP1074617-A2.
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AC ACHIEVE 2001: 2000EP-0116126.
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BP 27-AUC-1959; 95JP-0246036.
XX
AC 25-JUL-1959; 95JP-0200233.
AC 25-JUL-1959; 95JP-0300233.
AC 25-JUL-1959; 95JP-0300233.
AC 25-JUL-1959; 95JP-0300233.
AC 25-JUL-1959; 95JP-0300253.
AC 25-JUL-1959; 95JP-03
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Best Local Similarity 100.0%; I
Matches 22; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human cDNA sequence SEQ ID NO:16978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isogai T, Nishikawa T, Hayashi K, Saito K, Yamamoto J;
, Sugiyama T, Wakamatsu A, Nagai K, Otsuki T;
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Pred. No. 10;
0; Mismatches (
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Primer sets for synthesizing polynucleotides, particularly the 3602 full-length cDNAs defined in the specification, and for the detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs.

WPI; 2001-318749/34.

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RESULT 43

ANOSS118

XX ANOSS1

XX ANOSS1

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Matches 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Staphylococcus epidermidis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Probe: S.aureus: S.epidermis: E.taecalis: K.pneumoniae: E.cloacae: clinical sample:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Staphylococcus epidermidis probe SE-22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ55138;
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WPI; 1994-035086/04
                                                                         Eda S, Matsuhisa A,
                                                                                                                                                  (FUSO ) FUSO PHARM IND LTD (OHNO/) OHNO T.
                                                                                                                                                                                                                                                                  07-JUL-1992;
                                                                                                                                                                                                                                                                                                                                 07-JUL-1993;
                                                                                                                                                                                                                                                                                                                                                                                                          20-JAN-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9401583-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ55138 standard; DNA; 8654 6P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             al Similarity 100.0%;
22; Conservative U
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                                                                                                                                                                                                                                                                                                                                 93WO-JP00936
                                                                             Ohno T,
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v: Mismatches
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ds.
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Query Match
Best Local Similarity 100.0%; P
Matches 22; Conservative 0;
                                                                                                                                 The nuclectide sequence of an 8654 Lp probe obtained by digestion of Staphylococcus epidermials genomic DNA with the restriction enzyme Hindlll. The probes (AAQS5131-48 and AAQS5132-50) represent other probes derived, by Hindlll digestion of the genomes of Saureus, Spidermis, Enterococcus faccalis, Fseudomonas aeruginosa, E.coll, Slabisabla penumoniae or Enterobacter cloacae. The probes can be used to detect their respective microorganisms in clinical samples.
                                                                                                                                                                                                                                                                                                                                  Probe for identifying betteria causing infectious disease consists of a DNA fragment obtained by Hindill cleavage of pathogenic tacterial genomic DNA
                                                                                                                                                                                                                                                                                                Claim 4; Page 30-35; 100pp; Japanese.
                                                                                                Sequence 8654 BP; 2968 A; 1137 C; 1362 G; 3186 T; 1 other;
  Score 22; DB
; Pred. No. 8;
0; Mismatches
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     Gaps
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RESULT 4
AA162937
ID AA16
Husan genomic DNA SEQ ID NO 265.
                         22-GCT-2001 (first entry)
                                                                      AA162937 standard:
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                                                                      DNA; 11617
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11-JAN-2000; 2000US-0179065
01-FEB-2000; 2000US-0180028
15-HAY-2000; 2000US-025151
07-JUL-2000; 2000US-021680
11-JUL-2000; 2000US-021680
11-JUL-2000; 2000US-0225447
01-SEP-2000; 2000US-0230437
06-SEP-2000; 2000US-0230437
06-SEP-2000; 2000US-0230437
06-SEP-2000; 2000US-0231443
35-SEP-2000; 2000US-0231449
25-SEP-2000; 2000US-02314997
25-SEP-2000; 2000US-02314997
15-CCT-2000; 2000US-0234697
08-NOV-2000; 2000US-0246476
08-NOV-2000; 2000US-0246676
08-NOV-2000; 2000US-0246826
17-NOV-2000; 2000US-0246910
17-NOV-2000; 2000US-0249210 Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immunosuppressive; antiinflammatory; anti HIV antibacterial; vulnerary; antiparthismonian; antistekling; antianeanic; antiarthillic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiInflammatory; antiallergic; artidianetic; antipareatic; an 17-JAN-2001; 2001WO-US01346 02-AUG-2001. K0200155449-A1. Homo sapiens.

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RESULT
AAV3445
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AC AC

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to novel genes (AAI62752-AAI62961) and proteins (AAM2317-AAM42415) useful for preventing, treating or ameliarating conditions e.g. by protein or gene therapy. The genes are called conditions e.g. by protein or gene therapy. The genes are called from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful cand ovarient cand other cancers of the adrenal gland, bone, bone and ovarient cand other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract. Ever, lung, or urogenital: (b) immune disorders e.g. Addison's disease, allergies, autoimmune the manufacture of the adventure of the adventure of the selection's colities; (c) cardiovascular disorders such as myocardial isothemas; (c) cardiovascular disorders such as myocardial isothemas; (c) disease, multiple sclerosis, rheumatoid arthritis and ulcerative colities; (c) cardiovascular disorders such as myocardial isothemas; (c) disease, in firetious diseases such as viral, bacterial, fungal content and parasitic infections. Infectious diseases such as viral, bacterial, fungal content passitic infections, but was obtained in electrooic format directly from Wife at fip.vipo.int/pub/published_pcr_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Matches 22
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                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; SEQ ID NO 265; 532pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel plasma membrane associated proteins useful for disprasing treating, preventing and/or prognosing disorders related to the proteins, including cancer, immune response and neuronal disorders
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                                                                                                                                                                                                                                                                AAV34455;
                                                                                                                                                                                                                                                                                                        AAV34455 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 11617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                              4.5
                                                                                                                                                                               MHC class I chain-related gene A (MICA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ch 2.2%; So l Similarity 100.0%; F 22; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0249214
2000US-0249255
2000US-0250160
2000US-02511030
2000US-0251198
2000US-0251198
2000US-0251479
2000US-0251499
2000US-0251990
2000US-0251990
                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3043 A; 2768 C; 2512 G; 2854 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ruben SM;
                                                                                                                                                                                                                                                                                                        11722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 41: DB
; Pred. No. 7:7
0: Mis…atches
                                                                                                                                                                                                                                                                                                        נוי
טי
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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W09819167-A2
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                   /*tay= i
fnumber= 4
8732.8865
/*tsy= j
fnumber= 5
8870.11420
/*tag= k
fnumber= 5
11421.11722
/*tay= 1
                                                                                                                        /*tag= g
/number= 3
8354.8632
/*tag= h
/number= 4
8633.8731
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/number* 1
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110..6949
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/*Lag* c
/number* 1
6950..7204
/*Lag* d
                                                                                                                                                                                 /*tag= f
/number= 3
7767..8353
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/number- 2
7479..7766
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/number- 2
7205..7478
                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                       *cuntains
                                                                                                                                                                                                                                                                                                                                       11.11.06.5
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DR WFI: 1995-27339/24.

DR P-PSLE: AAM60043.

XX Use of MHC-related molecules, MICA and MICE - as targets for the PI detection and treatment of cancers and for isclating specific T cell populations which can be used for immunicible pay.

Spies 1. Spies V;

(HUTC-) HUTCHINSON CANCER KES CEMI FRED

29-001-1997; 29-001-1996;

97W0-USZJ170. 96US-0629044.

Disclosure; Page 164-111; 125pp; English.

This is the numan cell stress regulated MHC class I chain-related gene A (MICA). It was obtained from sligle-stranded (MUI) and double-stranded (MUI) templates of mapped or randomly shot-yan subcload DNA fragments that were derived from cosmid MIA. The game is located 40 kb centromeric of HIA-B. It codes for a 385 amino acid polypeptide (see AAM6004). A chik clone (see AAV34456) coding for closely-related MICB (see AAW60041) is also provided MICA and MICB are specifically expressed at the lining of the gastrointestinal tract, the primary site of infectious attack, and a major target for complications arising from graft-versus host disease (GVHD). They are also expressed at the cell surface of

cc colon and other cancer cells. A claimed method of detecting a calcied are methods of purifying, enriching and expanding voeta 1 craimed are methods of purifying, enriching and expanding voeta 1 craimed are methods of purifying, enriching and expanding voeta 1 craimed are methods of purifying, enriching purified Troells. Cc method of increasing expression of HICA or HICE pulyepides, a cc method of increasing expression of HICA by providing a cell with cc an expression construct comprising a HICA coding region under the Cc control of a promoter active in entaryotic cells, and a transgenic Cc animal expression MICA and/or HICE. A therapeutic agent (e.g. animal expression MICA and/or HICE. A therapeutic agent (e.g. a toxin or cytokine) conjugated to a HICA or HICE-hidning agent CC (e.g. antibodies) can be used for the detection and treatment of C concers such as brain cancer, lymphatic cancer, leakamia. Cc combandoma, head and neck cancer, ossophageal cancer, colon cancer. Cc melanoma, head and neck cancer, oversian cancer and creatment of other disorders such as CVHD and inflammatory boxel colleges.

Sequence 11722 BP; 2414 A; 2996 C; 2828 G; 3484 T; 0 other;

Query Match 2.2%; Score 22; DB 19; Length 11722; Best Local Similarity 100.0%; Pred. No. 7.7; Matches 22; Conservative 0; Hismatches 0; Indels 0 0; Indels 0; Gaps

0:

Search completed: April 30, 2002, 11:14:24 Job time: 12133 sec

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Title:
Perfect score:
Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                             Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                   Post-processing: Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OM nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      on:
                                                                                                                                  Pred. No. is the number of results predicted by chance to lawe a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic search, using sw model
                                                                                                                                                                                                                    l: //sqn2_6/ptodata/l/ina/5a_COMB.seq:*
l: //sqn2_6/ptodata/l/ina/5B_COMB.seq:*
2: //sqn2_6/ptodata/l/ina/6B_COMB.seq:*
3: //sqn2_6/ptodata/l/ina/6B_COMB.seq:*
4: //sqn2_6/ptodata/l/ina/6B_COMB.seq:*
5: //sqn2_6/ptodata/l/ina/backfiles1.seq:*
6: //sqn2_6/ptodata/l/ina/backfiles1.seq:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OLIGO_NUC
Gapop 60.0 , Gapext 60.0
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575
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(Without alignments)
422.095 Million cell updates/scc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gencore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                     SUMMARIES
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Description
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## 1 US : 09 - 04 - 054 - 27 US : 07 - 94 - 457 - 1 US : 08 - 43 - 32 - 1 US : 08 - 13 - 32 - 1 US : 08 - 13 - 32 - 1 US : 08 - 13 - 62 - 1 US : 08 - 13 - 565 - 14 US : 08 - 13 - 655 - 14 US : 08 - 13 - 655 - 14 US : 08 - 13 - 655 - 14 US : 08 - 13 - 655 - 12 US : 08 - 13 - 655 - 12 US : 08 - 13 - 655 - 12 US : 08 - 13 - 655 - 12 US : 08 - 15 - 655 - 2 US : 08 - 15 - 655 - 2 US : 08 - 15 - 556 - 3 US : 08 - 15 - 576 - 3 US : 08 - 16 - 576 - 3 Sequence 27, Applisequence 1, Applisequence 2, Applisequence 11, Applisequence 11, Applisequence 12, Applisequence 14, Applisequence 14, Applisequence 14, Applisequence 15, Applisequence 16, Applisequence 17, Applisequence 17, Applisequence 18, Applisequence 18, Applisequence 19, Applisequence 27, Applisequence 27, Applisequence 28, Applisequence 37, A Overy Match 55.5%, Score 31%; Db 4; Length 65. Best Local Similarity 100.0%, Pred No. 2.7e-129; Matches 315; Conservative 0; Mismatches 0; Indeis

Length 654;

0; Gaps

45	#	<b>.</b>	42	4 1	40	39	38	37	36	u.	4	نیا	32	31	36	25	
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US-09-032-523-5	US-08-891-581-2	US-08-721-690-2	US-08-162-247-1	US-08-380-916-2	US-09-246-335-55	PCT-US\$5-06846A-133	PCT-US94-05083C-129	US-08-470-339-133	US-08-734-664A-133	US-68-735-021-133	US-06-470-335-133	US-06-341-016-1	US-08-469-660-133	US-08-734-591A-133	US-08-469-526A-133	US-08-249-322A-133	00 00 403 303 433
Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	nedactice
5, Appl	<ol><li>Appli</li></ol>	<ol><li>Appli</li></ol>	<ol> <li>Appli</li> </ol>	2, Appli	55, Appl	133, App	•	133, App	133, App	133, App	133, App	<ol> <li>Appli</li> </ol>	133, App	133, App	133, App	133, App	****

ALIGNMENTS

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: INFORMATION FOR SEO ID NO. 2
SEQUENCE CHRACITEXISTICS:
SEQUENCE CHRACITEXISTICS:
TYPE ILLUCTUATION OF THE TOPOLOGY:
TOPOLOGY: 1 Inear
US-04-040-544-27
                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: SEED and BERKY LLP

STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
STATE: MA
COUNTRY: USA
ZIP: 56104
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKette
COMPUTER: IBM COmpatible
OPERATING SYSTEM: DS
SOFTMARE: FORSIEM: DS
SOFTMARE: FORSIEM: DS
SOFTMARE: FORSIEM: US/09/040,964
FILING DATE: 11 MAR-1998
CLASSIFFCATION:
ARPHICATION NUMBER: US/09/040,964
FILING DATE: 18 MAR-1998
                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: MAKI, DAVID J.
RECISIARION NUMBER: 31,352
RECERENCE/DOCKET NUMBER: 210121.456
ILLEDOMUNICATION INFORMATION:
TELEPHONE: 206-22-4500
TELEPHONE: 206-22-6031
TELEPHONE: 206-22-6031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          quence 27. Application US/09040984
Lent No. 6210883
ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANI: Reed, Steven G.
APPLICANI: Wang, Ingforg
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR DIAGNOSIS
TITLE OF INVENTION: OF LUNG CANCER
NUMBER OF SEQUENCES: 86
CORRESPONJENCE ALDRESS:
                                                                                                                                                27:
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b 148 CTAMARCHINTOCTIANTOCTIANTOCATAROAMACCTITICALITICAL HIGH HILLIH HIL
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| CONTINUE | 120 | Continue | March | 100 A; Score | 1, 100 A; Sco
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Covery Match 4.0%; Score 23; Lb1; Length 3207; Best Local Similarity 100.0%, Prod. No. 0.21; Length 3207; Matches 23; Conservative 0; Mismatches 0; Indels

0; Gaps

0

```
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/946,497
FILING DATE: 19921109
ATTONEY/AGENT INFORMATION:
AMBE: GENT: Stephen A.
REGISTRATION NUMBER: 29,768
REGISTRATION NUMBER: 16915/115
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5399
TELEPAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.H., Suite 500
GITT: Mashington, D.C.
COUNTR: USA
ZIP: 20017-5109
CAPPUTER REARABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC COMPUTER:
COMPUTER: LBM PC COMPUTER:
OPERATING SYSTEM: PC-POS/NAS-DOS
SOFTWARE: Patentin Release 11.0, Version 11.25
GURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/478,682
                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery March 4.0%; Score 3; DB 2; Length 32v7; Best Local Similarity 100.0%; Prod. No. 0.21; Marches 23; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                   08-520-678A-22/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEC ID NO. 1:
SEQUENCE CHRACTERISTICS:
LENGTH: 3007 base pairs
TYPE: DOLLER CAID
SPRANDEDNESS: double
TOPOLICKY INFORMER:
IMPELIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: HERRLICH, PETER APPLICANT: HERRLICH, PETER APPLICANT: PONTA, Helmut APPLICANT: GUENTHERF, UTSAL APPLICANT: HATZKI, Siegified APPLICANT: HATZKI, Siegified APPLICANT: HERZL ACHIM TITLE OF INVENTION: VARIANI CD14 SURPACE PROTEINS, DHA APPLICANT: HERZL ACHIM TOD14 SURPACE PROTEINS, DHA TITLE OF INVENTION: AS WELL AS THEIR USE IN DIAGNOSIS AND INFRAPY NUMBER OF SOURCESS: ADDRESS:
                                                                                                                                                                                                                                                                                                                                     PPLICANT: KOlykhalov, Alexander A.
TTLE OP INVENTION: NOVEL 3' TERMINAL SEQUENCE OF HEFATITIS
(TLE OP INVENTION: C VIRUS GENOME AND DIAGNOSTIC AND THEKAPEUTIC USES THEREOF
HRER OF SEQUENCES: 39
SHRESPONDENCE ADDRESS:
                                                                                                                                                                                  2, Application US/08520678A 5874565
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885575
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; TOPGLOGY: linear
; MCLECULE TYFE: DNA (genomic)
US-08-520-678A-22
                                                                                                                                                                                                                      INFORMATION FOR SEV ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGIH: 356 base pairs
TYPE: Lucleic acid
STRANDELNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING LATE:
CLASSIFICATION: 536
ATTONEZ/AGENT INFOGMATION:
ATTONEZ/AGENT I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIPE 63105
ZIPE 63105
ZICHOUTER, READABLE FORM:
REDURN TYPE; FICPLY disk
COMPUTER: LIFE, FICPLY disk
COMPUTER: STEM FOR COMPATILLE
COMPUTER: STEM FOR COMPATILLE
OPERATING SYSTEM: DECMOS/MS-LOS
SOFTMARE: PATENTION DATA:
APPLICATION NUMBER: US/08/520,678A
APPLICATION NUMBER: US/08/520,678A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACCEPSSEE: Howell & Haterker
STREET: 7733 Forsyth Blvd.,
CITY: St. Louis
STATE: MO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6029-6836
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```
CORRESPONDENCE ADDRESS:

ADDRESSEE: Howell & Haferkamp, L.C.
STREET: 7733 Forsyth Blvd., Suite 1400
CITY: St. Louis
STATE: MO
COUNTRY: USA
ZIP: 63105
COMPUTER KEADAGLE FUGM:
KEDIHN FREE Floppy disk
COMPUTER: 1LM FC Compatible
COMPUTER: FLM FC COMPATIBLE
COMPUTER: FLM FC COMPATIBLE
COMPUTER: FLM FC COMPATIBLE
COMPUTER: 1 MA FC COMPATIBLE
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COMPATIBLE
FILING DATE
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Best Local Similarity 100.0% Pred, No. 0.66;
Matches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3-08-597-126-22/c
Sequence 22, App
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2, Application US/08897126
6297003
                                                                                                                                                                                                                                                                                                                                                                                                                                       VIRUS GENOME AND DIAGNOSTIC AND THERAPEUTIC USES THEREGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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CLASSIFICATION:
PRIOR APPLICATION NUMBER: 08/520,678
FILING DATE:

EY/AGENT INFORMATION: Henderson, Melodie W.

0;

0

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HOLECULE TYPE: DNA (genomic)
US-08-897-126-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 3.8%; Score 22: DB 4: Length 355; Best Local Similarity 100.0%; Pred. No. 0.66; Matches 22: Conservative 0: Mismatches 0: Indeis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: February
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 37,848
REFERENCE/DOCKET NUMBER: 60:
LECOMMUNICATION INFORMATION:
TELEPHONE: 314-727-6188
TELEFAX: 314-727-6092
                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: doubl
TOPOLOGY: linear
DLECULE TYPE: cDNA
POTHETICAL: NO
                                                                                                                                                                                                                                                                     AME: KIRKPATRICK, ANITA M.
EGISTRATION NUMBER: 32,617
EFERENCE/DOCKET NUMBER: NIF
ECOMMUNICATION INFORMATION:
ELEPHONE: 619-235-8550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LE OF INVENTION: ISCLATION AND CHARACTERIZATION OF A
LE OF INVENTION: NOVEL CHAPERONE PROTEIN
BER OF SEQUENCES: 26
ESPONDENCE ADDRESS:
ESPONDENCE ADDRESS:
DRESSEE: KNOBBE, WARTENS, OLSON & BEAR
REET: 620 NEWPORT CENTER DRIVE, SIXTEENIH FLOOR
ITT: NEWPORT BEACH
ATE: CA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NGTH: 356 base pairs
PE: nucleic acid
RANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ION FOR SEQ ID NO: 22:
CE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                    TFICATION:
                                                                                                                                                                                                                                AX: 619-235-0176
ON FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/08203505B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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1..1730
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                                                                                                                                                                                                                                                                                                        NIH089.001A
US-08-468-245A-4

US-08-468-245A-4

: Sequence 4, Application US/08468245A

: Patent NJ 5886148

: GENERAL INFORMATION:
                                                                                                                          uvery Match 3.6%; Score 22; DB 1; Length 1731; Best Local Similarity 160.0%; Pred. No. 0.5%; Matches 22; Conservative 0; Mismatches 0; Indels
```

```
RECEISTATION NUMBER: 30.162
REFERENCE/DOCKET NUMBER: 00786/071002
TELECOMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-5070
TELEFAX: 2015 1 NO: 4:
SEQUENCE CHARACIERISTICS:
LENGTH: 2010
LENGTH: 2010
TYPE: nucleic acid
STRANGEDNESS: double
105-07-664-475A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IB P5/2 Model 502 or 555X
OPERATING SYSIEM: MS-DOS (Version 5.0)
SOFIWARE: MODEL (Version 5.1)
CURRENT APPLICATION NOTA:
APPLICATION NUMBER: US/07/864,475A
FILING DATE: 04-06-1992
CLASSIFICATION: 435
PRIOR APPLICATION: 435
Ouery Match 3.8%; Score 22; LbB 1; Length 2010; Best Local Similarity 100.0%; Pred. No. 0.58; Matches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: U.S.A.
21F: 02110-2804
COMPUTER READABLE FORM:
MEDIUM IVER: 3.5° Diskette, 1
COMPUTER: 13 H P5/2 McAal FOR
                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/681,702
FILING DATE: 05-04-1991
AITORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
NAME: Paul T. Clark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LICANT: Schipani, Ernestina
ILE OF INVENTION: PAKATHYROID HORMUNE KECEPTÜK
ILE OF INVENTION: AND DNA ENCODING SAME
GBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pplication US/07864475A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Fish & Richardson
225 Franklin Street
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0; Gaps

ESSEE: Fish & Richardson P.C. ET: 225 Franklin Street

PARATHYROID HORMONE RECEPTOR AND LWA ENCODING SAME

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COMPUTER INDUSTRIES

CONTROLL

CORRETTE: STR PF COMPATIBLE

OREGITES: STR
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: NAME/KEY: TATA_Signal
: LCCATION: (2458)...(2462)
US-09-136-605-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 14
LENGTH: 8056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Overy Match

3.8%; Score 22: LB 3;

Best Local Similarity 100.0%; Pred. No. 0.53;

Matches 22; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
GRGANISM: Homo sapiens
                                                                                                                                                                                     AITORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                        LATE: September 15, 1994
CLASSIFICATION: 514
AFPLICATION DATA:
AFPLICATION NUMBER:
FILING NO.
                                                                                                                                                                                                                                                                                                                                                                                       WORDER WORD WAS LAND WAS LASSIFICATION TO THE WORD WAS LASSIFICATION TO THE WORD WAS LAND WAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ALDRESSEE: Seidel, Gonda, Levorgua & Konaco, P.C.
STREET: Two pean Center, Suite 1800
CITY: Philadelphia
STAIE: Pennsylvania
COMITRY: U.S.A.
NAME: MONACO, Daniel A.
REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 83
LECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LE OF INVENTION: ANTISENSE
LE OF INVENTION: OLIGONUCLECTIVES TARGETING COOPERATING ONCOGENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ICANT: Calabretta, Bruno
                                                     8321-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 8056;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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```

Ouer; March 93.8%; Score 22; DB 1; Length 8082; Best Lucal Similarity 100.0%; Pred. No. 0.53; Matrices 22; Conservative 0; Mismatches 0; Indels

0;

Gaps

ATTORNY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/COCKET NUMBER: 1113
TELECOMMUNICATION: TOROMATION:
TELEPHONE: 201-487-5600

1113-1-006

COMPUTER READABLE FORM:

MEDIUM TIPE: Flopy disk
COMPUTER: LIBE: Flopy disk
COMPUTER: LIBE: CEMPARIBLE
OPERATING SISTEM: PC-MOS/MS-IGS
SOFIMABE: CALERIT RELEASE \*1.0, Version \*1.30
CURRENT AFFLICATION MUMBER: MS-08/811.566
APPLICATION MUMBER: MS-08/811.566
FILMS DATE: 03-MAR-1957
CLASSIFICATION: 435
CLASS

STREET: FLOOR
STREET: FLOOR
CITY: Hackensack
STATE: New Jersey
USA

LE OF INVENTION: FUNCTIONAL DAW CLONE FOR HEPATITIS C LE OF INVENTION: VIRUS (HCV) AND USES THEREOF SER OF SEQUENCES: 21

David A. Jackson, Esq. 1 Hackensack Ave, Continental Plaza, 4th INFORMATION:

Charles et al

Application US/08811566

```
SIREL: 400 Garas. City Flaza

CINTY: USA

CANTON USA

COMPUTER NAME

COMPUTER SEADNEE FOM:

COMPUTER SEADNEE FOM:
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Wed May 1 07:51:20 2002

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Page 7
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TOPOLOGY: linear
HOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-B11-566-1
                                                                                    TOPOLOGY: linear
MOLECULE TYPE: CUNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-B11-566-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INTILICANT: Rice, Charles et al.
TITLE OF INVENTION: FUNCTIONAL DIA CLONE FOR HEPATITIS C
TITLE OF INVENTION: VIRUS (HCV) AND USES THEREOF
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson, Esq.
STREET: Floor
STREET: Hockensack Ave. Continental Plaza, 4th
CITY: Hockensack
STATE: New Jorsey
COUNTRY: USA
21P: Offin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 15
US-08-811-566-5/c
: Sequence 5, Application US/08811566
: Patent No. 6127116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 3.8%; Score 22; Best Local Similarity 100.0%; Pred. No. Matches 22; Conservative 0; Mismatc
Ouery Match 3.8%; Score 22; DB 3; Length 12590; Best Local Similarity 100.0%; Pred. No. 0.51; Natches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                        TELEFAX: 201-343-1684
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 12980 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TVEE: Floppy disk
COMPUTER: IBH PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/08/811.566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                        TTORNEY/AGENT INFORMATION:
NAME: JACKSON ESQ., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 1113
TELECOMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 201-343-1684
FORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9646 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     eic
                                                                                                                                                                                                                                                                                                                                                                                           03-MAR-1997
N: 435
                                                                                                                                                                                                                                                                                                                     1113-1-006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 3; Length 9646;
  0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
  0
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HS-08-153-0518-27
Ap
                                                                                                                                                                                                                                LENGTH: 157

TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-153-0518-27
                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 32,327
REFERENCE/CACKET NUMBER: 264/
ITELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 469-1600
ITELEFAX: (213) 555-0440
ITELEX: (67-3510
INFORMATION FOR ESD ID NO: 27:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5495 AAAAAGGAAAAAAAAAAAAAAA 5474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET 637 West Fif STREET 53 West Fif STREET 50116 4700 CHY: Los Angeles STATE: California COUNTRY: U.S.A. 21P: 95071
                                                                      SIRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08153051B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Calvin B. Harley
Scott L. Weinrich
Catherine Strahl
Michael J. McEachern
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Michael D. West
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , W. Shay
ring E. Wright
abeth Blackburn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                            204/195
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RESULT 17
US-08-060-552C-43
; Sequence 43, Application US/08060952C
; Patent No. 569532
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guery Match 3.7%; Score 21; DB 1; Best Local Similarity 100.0%; Fred. No. 1.5; Matches 21; Conservative 6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/05/153,0518
FILING DATE: No. 5645986ember 12, 1993
PRIOR APPLICATION DATA: NO. 604/038,766
APPLICATION NUMBER: 08/038,766
FILING DATE: MATCH 24, 1993
ATTORNETY/AGENI INFORMATION:
NIMES VICTORIANION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER KENDABLE FORM:
MEDIUM TYPE: 3.5 "Diskette, 1.44 ML
MEDIUM TYPE: Storage
COMPUTER: SEM COMPALINE
COMPUTER: NEW COMPALINE
SOFTMARE: SENION: DEM PC DOS 5.0
SOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PEPLICANT: Michael J. McEachern
APPLICANT: HOMAYOUN VAZITI
TITLE OF INVENTION: THERRAPY AND DIAGNOSIS OF
TITLE OF INVENTION: CONDITIONS RELATED TO TELECMERE
TITLE OF INVENTION: LEGITH AND/OR TELOMERASE ACTIVITY
NUMBER OF SEQUENCES: 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 157;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
```

PEPLICANT: Michael D. West

APPLICANT: Jerry W. Shay

APPLICANT: Woodring E. Wright

APPLICANT: Elizabeth Eleckborn

APPLICANT: Elizabeth Eleckborn

ITLE OF INVENTION: RELATED TO TELOMERS

ITLE OF INVENTION: RELATED TO TELOMERS

TITLE OF INVENTION: RELATED TO TELOMERS

TOTLE OF INVENTION: RELATED TO TELOMERS

TOTLE OF INVENTION: RELATED TO TELOMERS

WHERE OF SEQUENCES: 57

```
STREET: 533 Was FIRTH Street

STREET: 512 Was FIRTH STREET

COMPUTER RANAGE FORM:

REDIEW TYPE: 3.5 Diskette. 1.44 Mb

MEDIEW TYPE: 3.5 Diskette. 1.42 Mb

MEDIEW TYPE: 3.5 Diskette. 1.44 Mb

MEDIEW TYPE: 3.5 Diskette. 1.42 Mb

MEDIEW TYPE: 3.42 Mb

MEDIEW
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Ouery Match 3.7%; Score 21; Lb Best Local Similarity 100.0%; Pred. No. 1.9 Matches 21; Conservative 0; Mismatches

2; Length 157;

Application US/08819867

I: HOMAYOUR WAZIFI
INVENTION: THERAPY AND DIAGNOSIS OF
INVENTION: CONDITIONS RELATED 10
INVENTION: TEOLOMERE LENGTH AND/OR

ELOMERASE ACTIVITY

Jerry Shay Woodring E. Wright Wirabeth H. Blackburn

herine M. Straht hael J. Mceachern

ωγοπ & Lyon 633 West Fifth Street Suite 4700

```
INILE OF INVENIEUR: CONMITTIONS BELEATED IN JIELDMERE

INTEE C INVENIEUR: LENTH AND/OR TELOMERASE ACTIVITY

KINDER CT SECULRICE SO

CORRESPONDANCE ALYON

STREET: SUITE 4700

CITY: LOS ANGRIES

COUNTRY: U.S. A.

COMPUTER REALABLE FORM:

MEDIUM TYPE: 3.5 UDISACTED SO

CONTRY: U.S. A.

COMPUTER REALABLE FORM:

MEDIUM TYPE: 3.5 UDISACTED SO

CONTRY: U.S. A.

COMPUTER REALABLE FORM:

MEDIUM TYPE: 3.5 UDISACTED SO

CONTRACE: LOS ANGRIES

COUNTRY: U.S. A.

COMPUTER: JEH COAPABILIE

OPERATION SYSTEM: INPO.

CORPORATED ANGRIES

COUNTRY: U.S. A.

APPLICATION NUMBER: U.S.

CORPORATION DATA: A.

APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FROR APPLICATION NUMBER: 08008,766

FROR APPLICATION NUMBER: 08008,766

FILLING DATE: NO. 5800644ember 12, 1953

FROR APPLICATION NUMBER: 08008,766

FROR APPLICATION NUMBER: 08008,7
```

CITY: Los Angeles STATE: Callfornia COUNTRY: U.S.A. ZIP: 90071-2066

```
NAME: Chambers Danial M. REGERRATION UNBER 31,561
REFERENCE/DOCKET NUMBER 324/5
TELEPHONE: C113) 469-1600
TELEPHONE: C113) 469-1600
TELEFAS: (213) 469-1600
TELEFAS: (213) 459-1600
TELEFAS: (57-3510
INFORMATION FOR SEC ID NO: 57:
SEQUENCE CHARCTERISTICS:
LENGTH: 156 base pairs
TYPE: INCIDENT acid
STRANDENESS: SINGLE
TOPOLICAY: linear
TOPOLICAY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER EXDABLE FORM:

MEDIUM TYPE: 3.5 DISKETTE, 1.44 Mb

MEDIUM TYPE: STOTAGE
COMPUTER: IBM COMPATIBLE
COMPUTER: BESTOTAGE
COMPUTER: BESTOTAGE
COMPUTER: BESTOTAGE
COMPUTER: BESTOTAGE
COMPUTER: BESTOTAGE
COURSENT APPLICATION DATA:
APPLICATION WHEER: US/08/815,867
FILING DATE: MATCH 14, 1957
FILING DATE: MATCH 14, 1957
FILING LASSIFICATION UNMERR: 08/15.051
FILING DATE: MO. 6007985ember 12, 1953
APPLICATION WUMBER:
TILING DATE: MO. FORTAGE
FILING DATE: MO. FORTAGE
APPLICATION WUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 3.7%; Score 21; DB Best Local Similarity 100.0%; Pred. No. 1.: Matches 21; Conservative 0; Mismatches
                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE
COMPUTER: IEN COMPALIBLE
OPERATING SYSTEM: DOS
SOFTMARE: FRSESSO for Windows Version 2.0
APPLICATION DATA:
FILING DATE: 18 MAR-1998
FILING DATE: 18 MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY (LP
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
STATE: WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT: Reed, Steven G.

PPLICANT: Wang, TongTong

ITTLE OF INVENTION: COMPOUNDS AND METHODS FOR DIAGH_SIS

TITLE OF INVENTION: OF LUNG CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2, Application US/09040984
6210883
(/AGENT INFORMATION:
Maki, David J.
                                                                                                                                                                                                                                                                                                      USA
                                                                      UMBER: US/09/010,984
18-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        224/232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Intels
```

ADDRESSLE: SIERNE RESSLEK, GGLDSIEIN & FOX, P.L.L.C. STREET: 1100 NEW YORK AVENUE, SUITE 600 CITY: WASHINGTON STAIE. DC.

STAIE: DC
COUNTRY: US
ZIP: 26065-5334
CAMPLIER RELABLE FORM:
KEDIUM TYPE: Floppy disk
COMPUTER: IMP Compatible
OPERATING SYSTEM: PC-DOS/MS-COS
SOFTHARE: PARCHITI RELEASE \*1.0, Version \*1.30
CAPPLICATION NUMBER: US/08/516.576B
FILING DATE:
CLASSIFICATION NUMBER: US/08/516.576B
FILING DATE:
APPLICATION NUMBER: US/08/516.576B
FILING DATE: S-36
PAIGA APPLICATION DATA:
CAPPLICATION NUMBER: US/08/516.576B
FILING DATE: 23-AUG-156-6
ATTORNEY/AGENT INFORMATION:
NAME: STEFFE ERIC K.
REDESERNE/COCET NUMBER: 1468.05000G1
TELECOMUMICATION NUMBER: 1468.05000G1
TELECOMUMICATION NUMBER: 1468.05000G1
TELECOMUMICATION NUMBER: 1500

FEATURE:
NAME/KEY: mat\_peptide

NAME/KEY: CDS LOCATION: 88..603

TOPOLOGY: linear MOLECULE TYPE: cDNA FEATURE:

TYPE: nucleic acid STRANDEDNESS: double

```
RESULT 21
US-0B-916-576B-3
Sequence 3, Application US/08916576B
Patent No. 6171816
                                                                                                                                                                                                                                                                                                                                                                                    Ouery Match. 3.7%; Score 21; Db 4; Length 217;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 21; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 21 ELECOMMUNICATION INFORMATION: 1ELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORMATION FOR SEQ ID NO: 82:
EQUENCE CHARACTERISTICS:
LENGTH: 217 base pairs
TYPE: nucleic acid
STRANGEDNESS: single
TOPOLOGY: linear
PPLICANI: ELHER, REINHARD
PPLICANI: EENER, REINHARD
PPLICANI: EIDERSS, GREGGRY A.
IILE OF INVENTION: NOVEL HUMAN GROWTH FACTORS
UMBER OF SEQUENCES 45
GRESPONIENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31, 394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         210121.456
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LOCATION: 88.156
US-08-916-576B-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-056-105-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100.0%; Pred. No. 1.6 Matches 21; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 157..603
FEATURE:
NAME/KEY: sig_pepti
                                                                                                                                                                                                                                                                                                                                                                                                          932 аааадуааааааааааааа 952
                                                                                                                                                                                                                                                                                                                                                                                                                           555 aaaaggaaaaaaaaaaa 575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LE OF INVENTION: VACCINES WITH ENHANCED INTRACELLULAR LE OF LIVENTION: PROCESSING REFERENCE: 233/221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 3.7%;
Local Similarity 100.0%;
les 21; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VT APPLICATION NUMBER: US/09/056,105
VT FILING DATE: 1998-04-06
PR APPLICATION NUMBER: 60/043,467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , Application US/09056105 6287569
                                                                                                                                                                                                                                                                                                                     , Application US/08437027
5670317
                                                                                                                                                                       2: Cooper & Dunham LLP
1185 Avenue of the Americas
                             PC compatible
M: PC-DOS/MS-DOS
ntin Release #1.0, Version #1.30
US/08/437,027
                                                                                                                                                                                                                                     DIAGNOSTIC TEST FOR TEST FOR THE DESMOPLASTIC MALL ROUND CELL TUMOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4; Length 1821;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ů; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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```
CLASSIFICATION: 536

ATIONERY/ABENT INFORMATION:

NAME: White John F

REDISTRATION HUMBER: 46416/JPH/CCA

FERENCE/DOCKET HUMBER: 46416/JPH/CCA

FERENCE/DOCKET HUMBER: 46416/JPH/CCA

FELEPHONE: 212 351 053

INFORMATION FOR 500 JD NO: 18:

SEQUENCE CHROCOTERTICS:

SEQUENCE CHROCOTERTICS:

LENGIH: 2412 base pairs

FIRE FORDINGS: single

SIRANGENESS: single

MOLECULE TYPE: CDNA to mRNA

US-08-437-027-18

Ouery Match

Best Local Similarity 100.0%; Pred. No. 1.6:

Best Local Similarity 100.0%; Pred. No. 1.6:
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ύγ 555 канаудаанандаандаа 575 ED 2391 ΑΛΑΚΩΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑ 2411

SESULT 24
US-66-1845-1
US-66-68703
Sequence 1. Application US/06891845
Sequence 1. Application US/06891845
Sequence 1. Application US/06891845
Patent No. 66568703
GENERAL INFORMATION: Gamma-Heregulin
APPLICANI SINEVENSION: Gamma-Heregulin
APPLICANI SINEVENSION: Gamma-Heregulin
UNDMER OF SOUNCESS 11
NUMBER OF SOUNCESS
APPLESSEE: Geneniech. Inc
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
CONHIN: OSA
ZIP: 4060
COMPUTER REALAGHE FOGM:
COMPUTER REALAGHE FOGM:
COMPUTER: 1848 C computible
OPERATING SVEITE: PC-COS/MS-OS
SOFTMARE: MINPATIO (Geneniech)
APPLICATION NUMBER: US/08/691.645
FILLNS DATE:
FILLNS DATE:
FILLNS DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/691.645
FILLNS DATE: HOROMATION:
REDEBERCE/CACKET NUMBER: 40.376
REDEBERCE/CACKET NUMBER: 40.376
REPERENCE/CACKET NUMBER: 40.376
REPUBLICANION: 40.476
REPUBLICANION: 40.476
REPUBLICANION: 40.476
REPUBLICANION: 40.476
REPUBLICANION: 40.476
REPUBLICANION: 40

Best Local Similarity 100.0%; Pred. No. 1. Matches 21; Conservative 0; Mismatches

```
TOPOLOGY: linear:
HOLECULE TYPE: other nucleic acid
US-08-619-5428-30
                                                                                                                                                                                                                                                                                                                                Query Match 3.5%;
Best Local Similarity 100.0%;
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/615,542B
FILING DATE: June 21, 1996
CLASSIFICATION: 194
ATTORNEY/AGENT INFORMATION:
NAME: WAILE, John P.
RECISTRATION NUMBER: 42840-A-PCT-US
REFERENCE/DOCKET NUMBER: 42840-A-PCT-US
TELEPOMUNICATION INFORMATION:
TELEPAX: (212) 391-0525
TELEPAX: (212) 391-0525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1185 Avenue of the Americas CITY: New York
STATE: New York
COUNTRY: USA
ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SED ID NO. 30:

SEQUENCE CHARACTERISTICS:

LENGTH: 339 base pairs

TYPE: nuclet acid

STRANDEDRESS: single

TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Pateentin Release #1.0, Version #1.3U
CUBRENT APPLICATION DATA:
                                                                                                                                                                                                                                                            PLE OF INVENTION: METHOD FOR CONSIRUCTION OF NURMALIZED THE OF INVENTION: CDNA LIBRARIES
AMT: Endege, Wilson O.
AMT: Steinmann, Kathleen E.
AMT: Astle, Jon H.
AMT: Burgess, Christopher C.
AMT: Bushnell, Steven E.
AMT: Catino, Theodore J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ), Application US/08619542b
5830662
                                                                                                                                                             pplication US/09328111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the Trustees of Columbia University in the City in the City New York
                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                Score 20; D6 2; Length 339;
Pred. No. 4.6;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 27
US-08-036-555B-133
                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM
OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
```

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; ORGANISM: Homo sapiens
US-09-328-111-25
                                                                                                                                                                                                                                                                  TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS
FILE REFERENCE: CCD-257 (UB)
CUNKENT APPLICATION UNMER US/05/32a,111
CUNKENT PILLING DATE: 1559-06-05
EARLIER APPLICATION UNMER US/05/32a,111
EARLIER APPLICATION UNMER US/05/32a,101
EARLIER APPLICATION UNMER US/05/32a,101
EARLIER FILMG DATE: 1559-06-10
```

Opery Match 3.5%; Score 20; DB 4; Length 413; Best Local Similarity 100.0%; Pred, Mo. 4.8; Matches 20; Conservative 0; Mismatches 0; Indels

e 133, Application US/08036555B No. 5530109 LL INFORMATION:

PELICANI: Geodearl, Andrew: Strowant, Faul;
PELICANI: Minghetti, Luiss; Materfield, Michael: Marchioni, Mark;
PELICANI: Clen, Maio So: Hiles, lan
IILE OF INVENTION: Feparation and Use
UMBER OF SEQUENCES: 184

ALDRESSEE: Felfe & Lynch
STREET: 605 Third Avenue
CITY: New York City
STATE: New York
COUNTRY: USA
ZIP: 10022
CLAPUTER READABLE FORM: 5.25 Inch, J60 kb Storage
CRADUTTY: TO Diskette, 5.25 Inch, J60 kb Storage

APPLICATION NUMBER: 07/06/36,555B
FILING DATE: 24 MAR 1953
CLASSIFICATION: 435
FILOR APPLICATION DATA:
APPLICATION NUMBER: 07/965.173
FILING DATE: 25-CCT-1952
FILOR APPLICATION NUMBER: 07/940.369
FILING DATE: 03-SEP-1952
FILOR APPLICATION NUMBER: 07/907,136
FILING DATE: 03-DATA: 30-DATA: 30

ATE: UD: NI... JICATION DATA: IION NUMBER: U.K. 91 07566.3 DATE: 10-APRIL-1991 PAPELI INFORMATION:

; STRANDEDNESS: single ; TOPOLOGY: linear US-08-036-5558-133

(212) 838-3884
OR SEQ ID NO:
ARACTERISTICS:

LUD 5250.4

Overy Match 3.5%; Score 20; DB 1; Length 744; Best Local Similarity 100.0%; Pred. No. 1.6; Matches 20; Conservative 0; Mismatches 0; Indels

3, Application US/08469569 5606032

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Goodearl, Andrew; Stroobant, Paul;
Hinghetli, Luisa: Waterfield, Michael; Marchioni, Mark;
Chen, Maio Su; Hiles, Ian
ENTION: Glial Mitogenic Factors, Their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cuery Match 3.5% Score 20; Lb 1; Length 744; Best Local Similarity 100.0%; Pred. Mc. 4.6; Matches 20; Conservative 0; Mismatches 0; Indels
CLLSSEP CHAILON UNMERS: 08/036.555

PRIOR APPLICATION UNMERS: 08/036.555

PRIOR APPLICATION UNMERS: 07/965.173

PRIOR APPLICATION UNMERS: 07/965.173

PRIOR APPLICATION UNMERS: 07/965.173

PRIOR APPLICATION UNMERS: 07/940.385

PILING DATE: 03.5EP-952

PRIOR APPLICATION UNMERS: 07/9707.136

PILING DATE: 03.5EP-952

PRIOR APPLICATION UNMERS: 07/907.136

PILING DATE: 03.5EP-952

PRIOR APPLICATION UNMERS: 07/907.136

PILING DATE: 03.5EP-1952

PRIOR APPLICATION UNMERS: 07/907.136

PRIOR APPLICATION UNMERS: 07/963.703

APPLICATION UNMERS: 07/963.703
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OPERATING SYSTEM: PC-DOS

SOFTHARE: MORIDBERIECT

CURENT APPLICATION DATA:

APPLICATION NUMBER: US/08/249,322A

FILING DATE: 26-MAY-1994

OF LASSIFICATION: A78-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:

ADDRESSEE: Pelfe 4 Lynch
STREET: 805 Third Avenue
CITY: New York
CONTRY: NEW York
CONTRY: NEW YOR
ACOUNTRY: UDA
PER 1002
PER 1002
PER 1002
PER 1002
PER 1003
P
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PLICANI: Minghetti, Luisa; Materfield, Michael; Marchioni, Mark;
PLICANI: Chen, Malo Su; Hiles, Ian
ILE OF INVENTION: Felal Mitogenic Factors, Their
ILE OF INVENTION: Felaparation and Use
GBER OF SEQUENCES: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      33, Application US/08249322A
5716930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
linear
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```

COMPUTER: IBH

COMPUTER: IBH

OPERATING SYSTEM

SOFTMARE: MOTOPERICC

CURRENT APPLICATION DATA:

APPLICATION UNMERE: US/08/465,565

FILING DATE: 06-UN-1995

CLASSIFICATION UNMER: US/08/465,565

PRIOR APPLICATION DATA:

APPLICATION UNMER: 07/965,73

FILING DATE: 24-WAR-1993

APPLICATION UNMER: 07/965,73

FILING DATE: 25-WAR-1993

APPLICATION UNMER: 07/940,389

FILING DATE: 03-SEP-1992

LING DATE: 03-SEP-1992

R APPLICATION NUMBER: 07/907,138

PLICATION NUMBER: 07/907,138

LING DATE: 30-JUN-1992

APPLICATION DATA:

APPLICATION DATA:

TON NUMBER: 34,266 2/DOCKET NUMBER: LUD 5250.4 CATION INFORMATION:

CONTISTING H.

MBER: U.K. 91 07566.3 R: 07/863,703 APRIL-1992 COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 5.25 inch, 160 kb storaye

ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
STATE: New York

eparation and Use

Wed May 1 07:51:20 2002

; TYPE: nucleic acid ; STRANDEDNESS: single ; TOPOLOGY: Linear US-08-249-322A-133

SEQUENCE CHARACTERISTICS: LENGTH: 744

Query Match 3.5%; Best Local Similarity 100.0%; Matches 20; Conservative (

DB i; Length 744.

```
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTMARE: FRANCED for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,526A
FILING DATE: 06 June 1995
CLASSIFICATION: 413
PRIOR APPLICATION DATA:
ATTORNEY/ACENT INFORMATION:

NAME: Bicker-Brady, Kristina
REGISTRATION NUMBER: 35,109
REFERENCE/DOCKET NUMBER: 0458
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
                                                                                                                       FILING DATE: 03-007-1592
APPLICATION NUMBER: 07/940, 385
FILING DATE: 03-5EP-1592
APPLICATION NUMBER: 07/940, 385
FILING DATE: 03-5EP-1592
APPLICATION NUMBER: 03-07/963, 703
FILING DATE: 03-APPLICATION NUMBER: 07/963, 703
APPLICATION NUMBER: 0 K. 91 07-664
THING DATE: 03-4PPLICATION NUMBER: 0 K. 91 07-664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MPUTER READABLE FORM
MEDIUM TYPE: Disket
                                                                                                                                                                                                                                                                                                     ICATION NUMBER: 08/0
NG DATE: 24-MAR-1993
ICATION NUMBER: 07/9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boston
: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08169526A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stroobant, Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PREPARATION AND USE
                                                                                                                                                                                                                                                                                                                                           08/036,555
                                                                                                                                     7863,703
1992
K. 91 07566.3
                                      04585/00200A
```

CCMPUTER READABLE FORM:

REDIUM IVEE: 3.5° biskette, 1.44 ML
COMPUTER: IAM COmpatible Pentium
COMPUTER: IAM COmpatible Pentium
SOPTWARE: RoadBeriest (Version 7.0)
COMMENT APPLICATION DATE: (Version 7.0)
APPLICATION NUMBER: 05/05/734,591A
FILING DATE: 22-CCT-996
CLASSIFICATION: 356

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/470,335
APPLICATION NUMBER: 08/036,555
PRIOR APPLICATION NUMBER: 08/036,555
FILING DATE: 03-MAR-1953
PRIOR APPLICATION NUMBER: 07/945,173
PRIOR APPLICATION NUMBER: 07/945,173
PRIOR APPLICATION NUMBER: 07/946,355
PRIOR APPLICATION NUMBER: 07/940,355
FILING DATE: 03-5EP-1952

APPLICATION NUMBER: 07/663,703
FILING DATE: 03.4FK-1992
FATOR APPLICATION NUMBER: NW 91 07566.3
FATILING DATE: 10.0.4FK-1991
ATTORNEY/GENT INFORMATION:

eker-Brady, Kristina TON NUMBER: 39,109 VDCCKET NUMBER: 04585/0u260P

PPLICATION DATA: CATION NUMBER: 07/507,136 NG DATE: 30-JUN-1552

```
; IUPOLUGY: linear
US-08-469-526A-133
                                                                                                                                                                                                                                                                                                                                                                                                                        Query March 3.5%; St
Best Local Similarity 100.0%; I
Matches 20; Conservative. 0;
                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
CIIY: boston
                                                                                                       OF INVENTION: GLIAL MITOGENIC FACTORS, THEIR OF INVENTION: PREPARATION AND USE R OF SEQUENCES: 187
: Massachusetts
RY: U.S.A.
02110
                                                                                                                                                                                                                                                                              Application US/08734591A 54220
                                                                                                                                                  Stroobant, Paul
Minghetti, Luisa
Waterfield, Michael
Hiles, Ian
Marchionni, Mark
Chen, Mario
                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 744;
                                                                                                                                                                                                                                                                                                                                                                                                                            0:
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US-00-469-600-133
US-00-469-600-133
US-00-469-600-133
US-elent Co. 133. Application US/08469600
Sequence 133. Application US/08469600
Sequence 133. Application US/08469600
APPLICANT: Grynne, David I.: Marchionni, Mark;
APPLICANT: Grynne, David I.: Marchionni, Mark;
APPLICANT: MEBURNY, Robert N.
TITLE OF INVENTION: THEIR PREPARATION AND USE
UNUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS;
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
CTTY: BOSLON TYPE: DISNette, 5.25 inch, 360 kb storage
COMPUTER REJABLE FORM:
ENDING TYPE: DISNette, 5.25 inch, 360 kb storage
COMPUTER: HEBER PREPARATION AND USE
COMPUTER: STATE: MASSACHUSELE PROMISE
COMPUTER: BOSLON DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US/08/465.660
FILING DATE: 29-JAN-1953
FILING DATE: 10-DEC-1952
FILING DATE: 25-EP-1952
FILING DATE: 10-DEC-1952
ATTORNEY/ACENT INFORMATION:
NAME: CLARK, Paul T.
REGISTRATION NUMBER: 07/927, 337
FILING DATE: 10-DEC-1952
ATTORNEY/ACENT INFORMATION:
NAME: CLARK, Paul T.
REFERENCE/DOCKET NUMBER: 07/927, 337
FILING DATE: 10-DEC-1952
ATTORNEY/ACENT INFORMATION:
NAME: CLARK, Paul T.
REGISTRATION NUMBER: 07/927, 337
FILING DATE: 10-DEC-1952
ATTORNEY/ACENT INFORMATION:
NAME: CLARK, Paul T.
REGISTRATION NUMBER: 07/927, 337
FILING DATE: 10-DEC-1952
ATTORNEY/ACENT INFORMATION:
NAME: CLARK, Paul T.
REGISTRATION NUMBER: 07/927, 337
FILING DATE: 10-DEC-1952
ATTORNEY/ACENT INFORMATION:
NAME: CLARK, Paul T.
REGISTRATION NUMBER: 07/927, 337
FILING DATE: 07
```

overy Match 3 3.5%; Score 20; DB 3; Length 744.
Best Local Similarity 100.0%; Pred, Mc. 4.6;
Matches 20; Conservative 0; Mismatches 0; Indels

0; Gaps

Application US/08470335F

Query Match 3.5%; Score 20; DB 2; Length 744; Best Local Similarity 100.0%; Pred. No. 4.6;

NAME/KEY: CDS LOCATION: (8)...(625) JS-08-470-335-133

CRGANISM: Bos taurus

FastSEQ for Windows Version 4.0

NUMBER: US/08/470,335f 1995-06-06 NUMBER: 08/036,555 NEG-03-24

CANI: HILES, IAN
OF INVENTION: GLIAL MITOGENIC FACTORS, THEIR
OF INVENTION: PREPARATION AND USE
REFERENCE: 04585/00200B

Usery Match 3.3%; Score 20; DB 2; Length 744; Best Local Similarity 100.0%; Pred, No. 4.6; Matches 20; Conservative 0; Hismatches 0; Indels

Wed May 1 07:51:20 2002

SEQUENCE CHARACTERISTICS:

133:

YPE: nucleic acid TRANDEDNESS: single >POLOGY: linear

```
EXRILER APPLICATION NUMBER: 07/863,703
EARLIER FILTING DATE: 1932-04-03
NUMBER OF SEG ID NOS: 192-04-03
SEOFTWARE: PASKSEQ FOR WINDOWS VERSION 3.0
SEOTT NO 13
LENGTH: 744
TYPE: DNA
TOPE: DNA
TOPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; NAME/KEY: CDS
; LOCATION: (8)...(625)
US-08-735-021-133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.5%;
Best Local Similarity 100.0%;
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                              OF INVENTION: GLIAL MITOGENIC FACTORS, THEIR OF INVENTION: PREPARATION AND USE REFERENCE: 04585/00200L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             al Similarity 100.0%;
20; Conservative (
        P: Stroobant, Paul
P: Minghetti, Luisa
P: Waterfield, Michael
Marchionni, Mark
P: Chen, Mario
P: Hiles, Ian
                                                                                                                                                                                                                                                                                                      Application US/08734664A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pplication US/08735021B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ION NUMBER: 08/036,555
ATE: 1993-03-24
TON NUMBER: 07/965,173
ATE: 1992-10-23
ION NUMBER: 07/940,389
ATE: 1992-09-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JMBER: 07/907.138
1992-06-30
IMBER: 07/863,703
592-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   995-06-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 3; Length 744;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGIH: 744
TYPE: nucleic acid
SIRANDEDRESS: single
TOPOLOGY: linear
US-08-734-664A-133
                                                                                                                                                                                                                                                                                                                                                                                                                       Ouery Match 3.5%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 4.6 Matches 20; Conservative 0; Mismatches
Sequence 133, Application US, Patent No. 6232366 GENERAL INFORMATION:
APPLICANI: GCODEARL, ANDREW APPLICANI: STROOBANT, PAUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION 536

FRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,322
FRIOR APPLICATION NUMBER: 08/249,322
FRIOR APPLICATION NUMBER: 08/36,555
FILING DATE: 24-ARK-193
FILING DATE: 24-ARK-193
PRIOR APPLICATION NUMBER: 07/565,173
FILING DATE: 23-CCT-1952
PRIOR APPLICATION NUMBER: 07/540,389
FILING DATE: 03-SEP-1952
PRIOR APPLICATION NUMBER: 07/540,389
FILING DATE: 03-SEP-1952
FRIOR APPLICATION NUMBER: 07/540,389
FILING DATE: 03-SEP-1952
FRIOR APPLICATION NUMBER: 07/540,389
FILING DATE: 03-JUN-1952
FRIOR APPLICATION NUMBER: 07/540,389
FILING DATE: 03-JUN-1952
FRIOR APPLICATION NUMBER: 07/540,389
FILING DATE: 03-JUN-1952
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MEDIUM TYPE: 3.5° Diskette, 1.44 ML
COMPUTER: 1BM COMPATIBLE PERLIUM
OPERATING SYSTEM: WINDOWS95
SOFTMARE: PastGey Version 2.0
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/06/734,664A
FILING DATE: 22 CGT-1956
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                              REFERENCE/COCKET NUMBER: 04565/002v0J
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 428-0200
TELEFAX: (617) 428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ITILE OF INVENTION: GLIAL MITGGENIC FACTORS, THEIR
TITLE OF INVENTION: PREPARATION AND USE
NUMBER OF SEÇUENCES: 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 176 Federal 5
CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE
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176 Federal Street
                                                                                                                             Application US/08470335C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JUN-1992
30-JUN-1992
ON DATA:
WHEEP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           UMBER: 07/863,703
03-APR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BER: UK 91 07566.3
0-AFK-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 744;
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NAME/KEY: CDS
: LOCATION: (8)...(625)
US-08-470-339-133
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Best Local Similarity 100.0%;
Matches 20; Conservative ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               S94-05083C-129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REENT APPLICATION NUMBER: US/08/470,335C
RRADAT FILING DATE: 1995-06-06
RLIER APPLICATION NUMBER: 03/036.555
RLIER FILING DATE: 1993-03-24
RLIER FILING DATE: 1993-03-24
RLIER FILING DATE: 1992-09-03
RLIER APPLICATION NUMBER: 07/907,138
RLIER FILING DATE: 1992-06-30
RLIER APPLICATION NUMBER: 07/663,703
RLIER APPLICATION NUMBER: 07/663,703
RLIER FILING DATE: 1992-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LE OF INVENTION: GLIAL MITOSENIC FACTORS, THEIR LE OF INVENTION: REPARATION AND USE REFERENCE: 04585/00208
                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: Massachusetts
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                           MPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANISM: Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LICANE: David I. Grynne
LE OF INVENTION: HETHODS FOR ALTERING
LE OF INVENTION: HUSCLE CONDITION
BER OF SEQUENCES: 185
RESPONDENCE ADDRESS:
                                                                                                                                                                                          PROTER: IBM
"WARE: Wordperfect
"WARE: Wordperfect
"WARE: Wordperfect
NAT APPLICATION DATA:
LICATION WARER: PCT/US94/05083C
LICATION WARER: PCT/US94/05083C
LICATION WARER: D6-MAY-94
                                                                                                                                                                                                                                                                                                                               UM TYPE: Diskette, 5.25 inch, 360
UM TYPE: kb storage
UTFR: THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     : Fish & Richardson
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application PC/TUS9405083C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Robert Sklar, Mark Marchionni,
NUMBER: 08/059,022
06-May-93
'INFORMATION:
                                                                                                                                               ON DATA:
                                                                                             MBER: 08/209,204
08-MAR-94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 744.
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SIRANDEDHESS:
TOPOLOGY: 11
PCT-US94-05063C-129
                         nucleic acid
    linear
                                          RACIERISTICS:
                                                                                                        30, 162
                                                                                               04565/028W01
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Odery Match 3.5%; S Best Local Similarity 100.0%; Matches 20; Conservative 0; 0;

US\$5-06846A-133 ence 133, Application PC/IUS9506846A ERAL INFORMATION: , Andrew David: Stroobent, Paul: i. Luiss; Materield, Michael; Marchioni, Mark; io Su; Hilles, Jan Glial Mitogenic Factors, Their

MFUTER KEADABLE FORM: MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage E: Felie & Lynch 805 Third Avenue Work City

COMPUTER: IBM
CFEMATING SYSTEM: PC-DOS
SOTTWARE: Wordperfect
CUKRON APPLICATION DATA:
APPLICATION NUMBER: PCI/U CT/US95/06846A

FILING DATE: 25-MAY-1995

FOR APPLICATION DATA:
APPLICATION UNDER: 08/249,322

FILING DATE: 26-MAY-1994

CLASSIFICATION:

APPLICATION NUMBER: 08/036,555

APPLICATION NUMBER: 08/036,555

FILING DATE: 21-MRR 1953

FRIOR APPLICATION NUMBER: 07/965,173

FILING DATE: 23-CCT-1952

FRIOR APPLICATION NUMBER: 07/940,369

FILING DATE: 03-SEP-1952

FILING APPLICATION NUMBER: 07/940,369

FILING APPLICATION NUMBER: 07/940,369

FILING APPLICATION NUMBER: 07/940,369

FILING APPLICATION NUMBER: 07/940,369

FILING DATE: 03-UNIA-1952

PRIOR APPLICATION NUMBER: 07/940,369 WATE: 30-JUN-1992
LICKTION DAIA:
IION NUMBER: 07/863,763
DAIE: 03-APRIL-1992
LICKTION DAIA:
LICK

: U.K. 91 07566.3 PRIL-1991

UD.5250.5

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PATENT NO. WASTERN TON:
APPLICANT. MCGONICLE. HRIAN
APPLICANT. MCGONICLE. BAIRN
APPLICANT. O'KEEF DANIEL
TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE ENZYMES
FILE REPERBNCE: CL-1126-A
CURRENT APPLICATION WHHEER: US/09/218,335
CURRENT FILING DATE: 1997-September-05
EARLIER APPLICATION WHHEER: OS/521,759
EARLIER APPLICATION WHHEER: OS/521,759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 3.5%;
Best Local Similarity 100.0%;
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 9 3.5%; Score 20; DB 5; Length 744;
Best Local Similarity 100.0%; Pred, No. 4.6;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
ZIP: 94111
APUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                         LOANT: Calydon, Inc.
E OF INVENTION: Tissue-Specific Enhancer Active 16.
E OF INVENTION: Prostrate
                                                                                                                                                                                                                                                                                                        Application US/08380916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/09248335
96504
                                                                                                                       Plehr, Hohbach, Test, Albritton & Herbert Eigharcardero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 20: DB 3; Length 934; Pred. No. 4.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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-162-247-1

, Application US/08182247 5830686

TILLE OF INVENTION: TISSUE-SPECIFIC EMHANCER ACTIVE IN
TITLE OF INVENTION: FROSTATE
NUMBER OF SEQUENCES: CONFESSOURCES ACRESCED LOADERSS:
ALDRESSEE: Loaneand and Townsend Kicurie and Crew
STREE: 1379 Lytich Avenue
STREE: California

CURFUER READABLE FORM:

#EDIUM TYPE: Floppy disk
CJAFOTER: IBM PC compatible
CFERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0.
CULFERI APPLICATION DATA:
CULFERI APPLICATION DATA:

Release #1.0, Version #1.25

US/08/182,247

16434-1

alifornia US

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MULICULE TYPE:
US-08-350-916-2
                                                               Query Match 3.5%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 4.4 Matches 20; Conservative 0; Mismatches
EFERENCE/DOCKET NUMBER: FP ECOMMUNICATION INFORMATION: ELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                  FIWARE: Patentin Rel
ENI AFPLICATION DATA:
PLICATION NUMBER: US
JUNG DATE: 12-JAN-19
                                                                                                                                                                                                                                                                                                                                                                      APPLICATION DATA:
ICATION NUMBER: US 08/182,247
NG DATE: 13-JAN-1954
EY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                : IBM PC compatible
33 SYSTEM: PC-DCS/MS-DOS
PARENTIN Release #1.0, Version #1.30
                                                                                                                                                            DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JAN-1995
                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                           FF-60058-FC
                                                                                               Length 1192;
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DNA (genomic)

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TELEX. 706141

TELEX. 706141

INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1192 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  overy Match 3.5%; score 20; DB 2; Length 1192; Best Local Similarity 100.0%; Pred: No. 4.4; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                    Query Match 3.5%; So Best Local Similarity 100.0%; F Matches 20; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Catherine, Polizzi M
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 34:
ELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PLICANT: Henderson, Daniel R.
TIE OF INVENTION: TISSUE-SPECIFIC ENHANCER ACTIVE
TIE OF INVENTION: IN PROSTATE
MEER OF SCOUENCES: 22
REESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ARE: PastSEQ for Windows Version 2.0 AppLICATION DATA:
"ATION NUMBER: US/08/77"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SSEE: MORRISON & FOERSTER
T: 755 PAGE MILL ROAD
Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/08721690 6057299
                                                                                                                                                                     TYPE: Genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                             ON NUMBER: US 08/380,916
TE: 30-JAN-1995
ON NUMBER: US 08/182,247
                                                                                                                                                                                                                                                                                                            415-494-0752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-SEP-1996
N: 514
                                                                                                                                                                                                                                                                                                                                                               34802-20001.21
                                                                    0; Indels 0;
                                                                                                                                                                                                                                                                                                                                          Owery Match 3.5%; Score 20; DB 3; Length 1192;
Best Local Similarity 160.0%; Pred: No. 4.4;
Matches 20; Conservative 0; Mismatches 0; Indels
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pplication US/09032523

0; Gaps

Incyte Pharmaceuticals, Inc.

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; MOLECULE TYPE: Genomic DNA
US-08-891-581-2
                                                                                                                                                                                                                          TELEM: 415-454-0/92
TELEM: 70611 NO: 2:
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
LENGTH: 1192 base pairs
TYPE: fucileic acid
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OPERATING SYSTEM: DOS

SOFTWARE: ESSÉSO for MINDOWS VERSION 2.0

APPLICATION NUMBER: US/08/891,561

APPLICATION NUMBER: US/08/891,561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Catherine, Polizzi M
REGISTIATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 34802-20001.22
TELECOMMUNICATION INFORMATION:
TELLEPHONE: 415-813-5600
TELEPAX: 415-415-690
TELEPAX: 415-41-694-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 514
PRION APPLICATION UNIA:
APPLICATION NUMBER: US 08/360,515
FILING DATE: 30-70x-1595
APPLICATION NUMBER: US 08/162,247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESPONDENCE ADDRESS:
LDRESSEE: MORRISON & FOERSTER
TREET: 755 FAGE MILL ROAD
TIY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/08691581
5136792
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ENTION: TISSUE SPECIFIC ENHANCER ACTIVE
ENLION: IN PROSTATE
THE PROPERTY OF THE PROPERTY 
                                                                                                                                                                                              single
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APPLICATION DATA: USASSEN TARES.
                                                                                  3174 Porter Drive
PE-0479 US
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Overy Match 3.5%; Score 20; DB 4; Length 1802; Best Local Similarity 100.0%; Pred: No. 4.3; Matches 20; Conservative 0; Mismatches 0; Indels 0; Indels 0; Gaps

Search completed: April 30, 2002, 10:54:09 Job time: 10918 sec

Page 19

endocrine; metabolism; regulation; melabsorption; gastritis; neoplesm.

Page

ARESULT 8
AAV34234 standard; D
ID AAV34234;
XX
AC AAV34234;
XX
C AAV34234;
XX
DE Human secreted prote
XX
Human; secreted prote
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XW
inmune system; asthm
XW
inflammation; ischae
XW
cognitive disorder;
XW
cognitive disorder;
XW
cognitive disorder; DNA; 1053 80

23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 21 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 26 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 20 - MAY 1597 21 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 26 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 21 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 21 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 21 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 20 - MAY 1597 21 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 20 - MAY 1597 21 - MAY 1597 21 - MAY 1597 22 - MAY 1597 23 - MAY 1597 24 - MAY 1597 25 - MAY 1597 27 - MAY 1597 28 - MAY 1597 29 - MAY 1597 29 - MAY 1597 20 - MAY 1597 20 - MAY 1597 20 - M

Human; secreted protein; tusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemis; developmental abnormality; foetal deficiency; blood; allergy; reani; Js; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS; cognitive disorder; senizophrenia; prostate; obestty; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;

Human secreted protein gene lo clone HMABH07.

28-JAN-1999 (first entry)

07-MAR-1997: 07-MAR-1997: 07-MAR-1997: mand saplets. 11-SEP-1998 WI 9639446-A2 \_C-MAR-1996; 58WC-0S04452

Sec 500 0+

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cc scretch duman protein. The gene number, and the clone it is derived from, are detailed in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene can be used to come it is compared to the human protein only. The gene can be used to come a compared to the human protein only. The first instance is acid sequences: ANV3145, 79476, amino acid sequences: ANV3154-79476, amino acid sequences ANV3155-795176; conditions can be disquested by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the completions of the new polypuches in a sample or by determining the amount of the completions in the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the amount of the new polypuches in a sample or by determining the new polypuches in the new polypuch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated human genes and the secreted polypeptides they encode useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-AUG-1997;
22-AUG-1997;
22-AUG-1997;
22-AUG-1997;
05-SEP-1997;
05-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 234; 447pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI: 1998-609887/51.
P-PSDB; AAW75137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bednarik DP, Brewer LA, Carter KC, Duan R, Ebner R, Endress GA;
Feng P, Perrie AM, Fischer CL, Graves KA, Greene JH, Hu JS;
Kyaw H, Lafleur DW, Li Y, Moore PA, Ni J, Olsen HS, Rosen CA;
Ruben SH, Shi Y, Soppet DR, Young PE, Yu GL, Zeng Z;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9705-0056882
9705-0056886
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Ouery Match 3.7%; Score 22; DB 19; Length 1053; Best Local Similarity 100.0%; Pred. No. 14; Matches 22; Conservative 0; Mismatches 0; Indels (

0; Indels 0; Gaps

0.

Sequence 1053 BP; 248 A; 298 C; 295 G; 209 T; 3 other:

GenCore version Copyright (c) 1993 - 2000

Compagen Ird

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Searched:
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1: /ogn2_6/ptcodate///ina/5A.COMB seq:*

2: /ogn2_6/ptcodate///ina/5B.COMB seq:*

3: /ogn2_6/ptcodate///ina/pcTUs_COMB seq:*

5: /ogn2_6/ptcodate///ina/pcTUs_COMB.seq:*

5: /ogn2_6/ptcodate///ina/pcTUs_COMB.seq:*

6: /ogn2_6/ptcodate//ina/pcTUs_COMB.seq:*
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Gapop 60.0 , Gapext 60.0
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(Githout allgoments)
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              1 US-08-833-45
2 US-08-832-87-19
4 US-08-870-115-10
3 US-08-187-115-17
3 US-08-188-787-72
2 US-08-858-787-22
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2 US-08-858-787-23
2 US-08-858-787-23
2 US-08-863-028-23
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1 US-08-872-85
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sequence 15, Apples equence 17, Apples equence 18, 
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4. Appr	Sequence	4 US-08-955-916C-4	338	1.9	18	<u>*</u>
14. App	Sequence	4 US-05-406-071-14	338	Ī	18	1
14, App	Sequence	3 US-09-248-203-14	338	•	18	43
14. App	Sequence	2 US-08-936-706A-14	338	•	18	<b>\$</b> 2
14. App	Sequence	2 US-08-536-707A-14	338	_	16	12
14. App	Sequence	2 US-08-922-267A-14	338	_	18	to
14. App	Sequence	1 US-08-825-553-14	338	_	16	36
14. Appl	Sequence	1 US-08-631-200-14	336	1.5	16	36
1, Appl	Sequence	4 US-05-073-257-1	333	_	16	37
37. App	Sequence	2 US-08-896-410-37	305		16	35
26. App	Sequence	1 US-08-684-672-26	854		16	u,
	Sequence	1 US-08-684-672-25	854	1.9	18	LJ da
	Sequence	4 US-09-461-474-13	750	1.9	16	w
	Sequence	4 US-09-328-111-572	621	1.9	16	
App		4 US-09-385-982-98	285	1.5	18	C 31
A, Appl	Sequence	1 US-08-578-645-4	561	1.9	16	30
I, Appl	Sequence	2 US-08-858-834-1	432	1.5	18	29
1. Appl.	Sequence	2 US-68-858-830-1	432	1.9	18	26

## ALIGNMENTS

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DESCRIPTION US/OB832883

DESCRIPTION US/OB832883

DESCRIPTION US/OB832883

DETECNATION BAIGHTON ANTONIO
APPLICAMIT BAIGHTON
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CORRESPECTORIACE ADDRESS:
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CORRESPECTORIACE ADDRESS:
CORRESPECTORIACE PARTICLE FORM:
APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: 30.480
REGISTATION ANTONION: 435
ANTE: MADRO DATA:
APPLICATION NUMBER: 30.480
REFERENCE/DOCKET NUMBER: 3
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Gaps

0

us-09-248-178-62.rni

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; MOLECULE TYPE: US-08-832-877-49
                                                                                                                                                                                                                                                                                                                                                                                                                                             opery Match 2.3%; Score 21: DB 2: Length 4576; Best Local Similarity 100.0%; Pred: No. 1.5; Natches 21: Conservative 0; Mismatches 0: Indois
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                                                                                                                                                                             EERAL INFORMATION:
PLICANT: BENDEE: MILSON O., ET AL.
TILE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TILE REFERENCE: CCDMA-250XX
                                                                                                                                                                                                                                                                                                                                                                           ESPONDENGE ADDRESS:
DDESSEE: SEIDEL, GOMDA, LAVORGNA & MONACO, P.C.
REET: Suite 1800 Two Penn Center Plaza
TY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ICANT: Giordano, Antonio

I INVENTION: METHODS FOR THE DIAGNOSIS AND PROGNOSIS OF
LE OF INVENTION: CANCER
LE OF SEQUENCES: 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ME: MODAGO, Daniel A.
GISTRATION NUMBER: 30,480
FERENCE/DOCKET NUMBER: 8321
COMMUNICATION INFORMATION:
EPHONE: (215) 568-8383
EFAV. (215) 568-8383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JTER READABLE FORM:
JUM TYPE: Floppy disk
HOTER: IBM PC compatible
ERATING SYSTEM: PC-DOS/MS-DOS
ETWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08832877
astSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                     oplication US/09385982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA (genomic)
                                                                                                                       ON NUMBER: US/09/385,982
(TE: 1999-08-30
ON NUMBER: 09/328,111
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                                                               WBER: 60/117,393
                          дея: 60/098,639
98-08-31
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-LOS
SOFTWARE: Patentin kelease 11.0, Version 11.30
CUREENT APPLICATION DATA:
APPLICATION NUMBER: US/08/870,126
FILING DATE:

RESPONDENCE ADDRESS:
DRESSEE: Howson and Howson
TREET: Spring House Corporate Cntr, P 0 Box 457
TTY: Spring House
TTY: Pennsylvania

INVENTION: Murine and Human Box-Dependent INVENTION: MYC-Interacting Protein (Bin1) and Uses Therefor

CLASSIFICATION: 530
PRIOR AFPLICATION DATA
APPLICATION HUMBER: US 08/435,454
FILING DATE: 05-MAY-1995
PAIGA AFPLICATION DATA:
AFPLICATION NUMBER: US 08/652,572
FILING DATE: 24-MAY-1956

NAME: KOGTOIF, Cathy A.
FETERSISHATION NUMBER: 33,580
LECOMMUNICATION INCOMATION:
LECOMMUNICATION INCOMATION:
LECTRONE: 215-540-9200

IELEFAX: 215-540-5818
BWATION FOR SEO ID NO:
DUENZE CHARACTEKISTICS:
DENGTH: 3226 base pairs
IYPE: nucleic acid
STRANGEDNESS: double

LOCATION: 851..945 OTHER INFORMATION:

/note\* \*exon 4\*

NAME/KEY: exon LCCATION: 677..734 OTHER INFORMATION: /note\* "exon 3"

ECULE TYPE: DNA (genomic)

NAME/KEY: exon
COCKTION: 24/3..2579
OTHER INFORMATION: /note\* "exon 6"
US-08-870-126-10

NAME/KEY: exon LOCATION: 1408..1503 OTHER INFORMATION: //

ouery Match 2.2%; Score 20; Db 3; Length 3226; Best Local Similarity 100.0%; Pred. No. 4.3; Matches 20; Conservative 0; Mismatches 0; Indels

0; Gaps

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0;

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Oy 262 tetecagagactrastics 281

Db 1297 TCTCAGACCCTCAGTTTCC 1278

RESULT 5

GUS-09-128-155-16/C

GUS-09-128-156/A

GUS-09-128-158/A

GUS-0
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Query Match 2.1%; Score 15; Dis 2; Length 28; Best Local Similarity 100.0%; Pred. No. 14; Matches 15; Conservative 0; Mismatches 0; Indels

0;

Gaps

506 aggcaaaaaaaaaaaaaaa 924

AGGCAAAAAAAAAAAAAA 25

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Query Match 2.1%; S
Best Local Similarity 100.0%;
Matches 19; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-005/MS-Dos
SOFTWARE: Parentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (202)672-5399
TELEX: 904136
FORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                   ASSIFICATION.

OR APPLICATION NUMBER: US 08/481.687

FILING DATE: 07-JUN-19-5

FILING DATE: 07-JUN-19-5

FICANEY/ACENT INFORMATION:
NAME: BBNT SCEPBEAN FRATION NUMBER: 29,768
REFERENCE/FOCKET NUMBER: 33229/325,
ELECOMMUNICATION INFORMATION:
FFI.SPHONE: (2023-57494)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ER OF SEDUENCES: 39
ESPONDENCE ADDRESS:
ESPONDENCE ADDRESS:
DRESSEE: FOLEY & Lardner
DRESSEE: 1000 K Street, N.W., Suite 500
REET: 1000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ICATION NUMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Washington
: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T: DUVICK, Jonathan P.
TIN BRIGGS, Steven P.
INVENTION: PCR-BASED CDMA SUBTRACTIVE CLONING INVENTION: MCTHOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      opplication US/08858767
                                                                                                                                         pplication US/08858767
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N: 435
PCR-BASED CDNA SUBTRACTIVE CLONING METHOD
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PPLICATION ...

PPLICATION ...

LASSIFICATION: 435

LASSIFICATION DATA:

APPLICATION MUMBER: US 08/481.687

FILING DATE: 07-9UN-195

FILING DATE: 07-9UN-195

FILING DATE: Stephen A., 78229/3.
                                                                                                                                                                   APPIER: Flogdy disk
APPIER: IBM PC comparible
SERATIKS SYSIEM: PC-DOS/MS-U/S
AFTMARE: PatentIn Release #1.0, Version #1.30
ENI APPLICATION DATA:
PLICATION MURREY.
                                                                                                                                                                                                                                                    HER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                   3300 K Street, N.W., Suite 500
  33229/325/PIHI
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Query Match 2.1%; Sco Best Local Similarity 100.0%; Pro-Matches 19; Conservative 0; 

Gaps

1, Application US/08863028 5853991

FICANI BUVICK, Jonathan P.
FICANI BRIGGS, Steven P.
TLE OF INVENION: PREMASED CDMA SUBIRACTIVE CLONING
TLE OF INVENION: METHOD
MAER OF SEQUENCES: 99
MAESFORIENCE ADDRESS:
ADDRESSEE: FOLEY & Lardner
MAESFORIENCE ADDRESS.

STATE: D.C. H TYPE: Ploppy disk
TER: IBM PC Compatible
TING SYSTEM: PC-DOS/MS-DOS
ARE: Patentin Release #1.0, Version #1.30
APPLICATION DATA: US/08/663,028

CATION: 435 ICATION DATA: IGN NUMBER: US/08/858,767 AIE: 19-MAY-1997

R: US 08/481,687 JUN-1995

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Query Match 2.1%; S. Best Local Similarity 100.0%; | Matches 19; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 904136

NFORMATION FOR SEQ ID NO: 21:
PRODIENCE CHARACTERISTICS:
                                                                                                                                                 PRIOR APPLICATION NATA:
APPLICATION NUMBER: US/08/858,767
FILING DATE: 19-MAY-1997
APPLICATION NUMBER: US 08/481,887
FILING DATE: 07-UN-1995
FILING DATE: 07-UN-1995
ANDRECY AGENT IMPORMATION:
ANDRECY AGENT IMPORMATION:
                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TIPE: Flopy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-POS/MS-DOS
SOFTMARE: PATEONITH Release *1.0, Version *1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/863,028
FILING DATE:
CLASSIETICATION: 435
CLASSIETICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 AGGCAAAAAAAAAAAAAA 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ME: BENT, Stephen A:
GISTRATION NUMBER: 29,768
FERENCE/DOCKET NUMBER: 3327
COMMUNICATION INFORMATION:
LEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                               : Washington
(: 904136
ION FOR SEQ ID NO: 22:
CE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2, Application US/08863028 5853991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DENCE ADDRESS:
SE: Foley & Lardner
SE: Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ICATION INFORMATION:
E: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                     33229/325/PIHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          33229/325/PIHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 28;
                                                                                Query Match
Best Local Similarity 100.0%; Pred. No. 14;
Matches 19; Conservative 0; Mismatches
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RESULT 13 US-09-248-335-29

Indels

0

Gaps

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US-08-653-028-22
ILLERON.

TELEX: 904336

INCAMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
TYPE: nucleic acid
"TARNEEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouer; Match 2.1%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 14 Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFAL INFORMATION AND AMERICANI PARKET STATE OF THE STATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCAPTIER READABLE FORM:

MELTUM TYPE: Floppy disk

COMPUTER: IBM PC compartible

COMPUTER: FLOM PC COMPARTIDE

COMPATING SYSTEM: PC-005/MS-106

SOFTWARE: Patentin Release #110, Version #1.36

SOFTWARE: Patentin Release #110, Version #1.36

CURRENT APPLICATION DATA:

105/04/664 008
                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRICE APPLICATION UNTA:
APPLICATION UNMER: US/08/658,767
FILING DAIE: 15-MAY-1997
APPLICATION UNMER: US 08/481,687
FILING DAIE: 07-JUN-195
AFILEATION UNMER: US 08/481,687
AFILEATION UNMER: 25,768
AFICATION UNMER: 25,768
AFICERNEY/COCKET UNMER: 31225/325/1
FILEDHONE: (202)672-5300
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Foley & Lardner SUITE: 3000 K Street, N.W., Suite 500 CITE: Kashington STAEE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/863,028 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3, Application US/08863028
5853991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       33225/325/PIHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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Wed May 1 07:51:05 2002

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Query Match 2.1%; Score 19; Best Local Similarity 100.0%; Pred. No. Matches 19; Conservative 0; Nismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: MICROSO
SEQ ID NO 29
SEQ ID NO 29
SEQ II 1134
TYPE: DNA
ORGANISM: maize
09-248-335-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-672-850-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: SILVA. RODIN M.
REGISTRATION NUMBER: 39.304
REFERENCE/DOCKET NUMBER: A-63
RELECOMMUNICATION NEFORMATION:
TELEPRONE: (415) 781-1989
TELEPRA: (415) 398-1398
ORMATION FOR SEG ID MO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE ENZYMES
                                                                                                                                                                                                                                                                                                                                                   San France:
E: California
MTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Microsoft Word Version 7.0A
                                                                                                                                                                                                                                                                TYPE: Floppy disk
RR: IBM PC compatible
RB: PC compatible
RB: Patentin RC:005/MS:003
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , Application US/09248335
6096504
                                                                                                                                                                                                                                                                                                                                         READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08672850
140117
                                                                                                                                                                                                                                                                                                                                                                                                  : Flehr, Hohbach, Test, Albritton & Herbert
Four Embarcadero Center, Suite 3400
n Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Milbrandt, Jeffrey
Araki, Toshiyuki
                                                            235 base pairs
leic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ION NUMBER: US/05/248,335
ATE: 1999-02-10
                                                                                                                                                                                                                                     24-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1997-September-05
                                                                                                                                                                                                                                                     US/08/672,850
                                                                                                                                                                A-63610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 1134;
```

Query Match 2.1%; Score 19; DE 3; Length 1235; Best Local Similarity 100.0%; Pred. No. 12;

Application US/08833623 5969093

Gaps

Jacobs, Kenneth McCoy, John Kelleher, Kerry

```
: NAME/KEY: CDS
; LOCATION: 38..1447
US-08-514-014-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 19; Conservative
Ouery Match. 2.1%; Score 19; Db 1; Length 2209;
Best Local Similarity 100.0%; Prod. No. 12;
Matches 19; Conservative 0; Mismatches 0; IndelB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MPDITER READABLE FORM:
MEDIUM TYPE: Floopy disk
MEDIUM TYPE: Floopy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTIWARE: Patentin Release #1.0, Version #1.25
SRENT APPLICATION DATA
APPLICATION NUMBER: US/08/514.014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DDRESSEE: Genetics Institute, Inc. -- Legal Affairs TIREET: 87 CambridgePark Drive ITY: Cambridge TATE: Massachusetts CONTRY: No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                      STRATION NUMBER: 32,724
STRATION NUMBER: 32,724
MRENCE/DOCKET NUMBER: GI
MANUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      upplication US/08514014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               arlin, McKeoug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA SEQUENCES AND SECRETED PROTEINS ENCODED THEREBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ċ.
```

ADDRESSEE: Genetics Institute, Inc. -- Legal Affairs STREET: #7 CambridgePark Drive CITY: Cambridge STATE: Massachusetts

EQUENCES AND SECRETED PROTEINS

```
: NAME/KEY: CDS
: LOCATION: 38..1447
US-08-833-823-1
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 2.1%; So
Best Local Similarity 100.0%; 1
Matches 19; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPACIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: PATENTION DATA:

APPLICATION NUMBER: US/08/933.823

FILING DATE: 10-74R-1957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELEFAX: (617) 876-5851
FORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 52,724
LECOMMUNICATION PARTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EY/AGENT INFORMATION:
                                                                                                                                                                                                                                                             pplication US/06813150
DNAX Research Institute
901 California Avenue
10 Alto
                                                                                                                                                                                           weller, Christopher
ebecque, Serge J.E.
                                                                                                                   ing, Lynette M.
ine, Constance M.
in, Daniel M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MBER: 08/514,014
                                                                                    WAMMALIAN PROTEINASES; OXIDOREDUCIASES;
                                                                                                                                                                                                                                                                                                                                                                                                                        Score 19; DB 2;
Pred. No. 12;
U; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 2205;
```

US95-09261-1

sence 1, Application PC/TUSy509261
NERAL INFORMATION:

BOARD OF RESENTS, THE UNIVERSITY OF TEXAS SYSTEM 201 West 7th Street Austin

CANI: POSTAL CODE: 78701
CANI: TELEPANE NO: C5121495-4462
ANT: TELEFAX: (512)499-4523
OF INVENTION: METHODS AND COMPOSITIONS FOR THE EXPRESSION OF DELIVENTION: A BONE AND PROSTATE DERIVED GROWTH FACTOR

United States of America E: 78701

E: Arnold, White a Durkee P.O. Box 4433

UNITED STATES OF AMERICA

COMPUTER READABLE FORM:

#EDIDIN TIPE: Floppy disk
COMPUTER: HAN PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-L/S/ASCII
SOFTWARE: PATENTION DATA:
APPLICATION NUMBER: PCT/USS5/05261
FILING DATE: CONCURRENTLY HEREWITH
FILING DATE: CONCURRENTLY HEREWITH

IOR APPLICATION DATA: APPLICATION NUMBER: US 08/283,701

```
: NAME/KEY: CDS
: LCCATION: 61..1470
US-08-613-150-1
                                                                                                    INCORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGIH: 2280 base pairs
IYFE: nucleic acid
STRANEERESS: single
IJOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                           HEDINI TYPE: Floppy disk
CCMPUTEK: IBM PC compatible
CPERATING SYSTEM: PC-TOS/MS-TOS
SUSTEMATE: Palestin Release 1.0, Version 1.30
CUBRENT APPLICATION DATA:
LEPTICATION NUMBER: UB/08/613.150
FILLNG DATE: COLORAN-1997
CILCUSTEMATE: COLORAN-1997
                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                  ITORNEY/AGENT INFO.....
KAME: CHING, EGWIN P.
REGISTRATION NUMBER: $1,090
REGERENCE/DOCKET NUMBER: $706
RELECOMUNICATION INFORMATION:
FELECOMUNICATION 1650-852-9196
                                                                                     LE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        S4304-1104
K READABLE FORM:
                                                                                                                                                                                                                                      650-496-1200
                                                                                                                                                                                                                                                                                                          SF0693
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vaery Match 2.1%; Score 19; DB 3; Length 2280; East Local Similarity 100.0%; Pred. No. 12; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                      0;
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FILING DATE: 01-AUG-1994 CLASSIFICATION:

INFORMATION:

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NAME/KEY: CDS;
; LOCATION: 694..2310
PCT-US95-09261-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match 5 initiatity 100.0%; Score 19; DB 5; Length 3171;
Best Local Similarity 100.0%; Prod. No. 12;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-612-521-3/c
NAME: Hamilton, Amy E
REGISTRATION NUMBER: 33,854
REFERENCE/DOCKET NUMBER: X-S
RELECOMMUNICATION INFORMATION:
TELEPHONE: 317-276-3169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 34,430
REFERENCE/DOCKET NUMBER: UTI
REFERENCE/TOCKET NUMBER: UTI
ELLECHIONE: (512) 418-3000
TELLECAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RMATION FOR SEQ ID NO: 1:
QUENCE: CHARACTERISTICS:
LENGIH: 3171 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                    RRESPONDENCE ADDRESS:

ADDRESSE: Patent Division/AEH
STREET: Lilly Corporate Center
CITY: Indianapolis
STATE: Indiana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LECULE TYPE: other nucleic acid
DESCHIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ), Application US/U8612521
), 5786463
                                                                                                                                                                                          LIK REMDABLE FORM:
IUM TYRE: Floopy disk
UUTRR: HEM POOPY disk
WITEN SYSTEM: PO-DOS/MS DOS
TUNKE: Potential Release 1.0, Version #1.30
                                                                                          HEY/AGENT INFORMATION:
                                                                                                                                       T APPLICATION DATA:
ICATION NUMBER: US/08/612,521
AG DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                      MULTIPLE DRUG RESISTANCE GENE OF CRYPTOCOCCUS NEOFORMANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UTFC422P--
                                       K-9693
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exon
1130..1395
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2913..3009
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1443..1872
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541..1072
exon
5553..5588
                     exon
5162..5500
                                          exon
5006..5113
                                                               excn
4793..4958
                                                                                     exon
4487..4743
                                                                                                           exon
4291..4438
                                                                                                                                 exon
4016..4244
                                                                                                                                                     exon
3831..3966
                                                                                                                                                                           exon
3614..3776
                                                                                                                                                                                                 exon
3438..3561
                                                                                                                                                                                                                       exon
3076..3379
                                                                                                                                                                                                                                                                  exon
2479...2863
                                                                                                                                                                                                                                                                                      exon
1925...2426
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RESULT 20
US-09-01-32A-50
US-09-01-32A-50
Sequence 50, Application US/0909132A
Patent Mo. 6284462
GENERAL INFORMATION:
APPLICANI: SUCA, TAKANUGI
APPLICANI: SUCA, TAKANUGI

OF INVENTION: PROBES AND METHODS FOR POLYNUCLEOTIDE DETECTION OF SEQUENCES: 58

Query Match 2.1%; Scot Best Local Similarity 100.0%; Promatches 19; Conservative 0; 1

0; Indels 0;

CORRESPONDENCE ADDRESS:
ADDRESSEE: LEYDIG, VOIT & MAYER, LID.
STREET: TWO PRUDENTIAL PLAZA, SUITE 4900
CITY: CHICAGO
STATE, II

```
CLASSIFICATION: 433
CLASSIFICATION BATA:
APPLICATION MUHBER: JP 962566:3
FILENG DAYE: JP 962566:3
FILENG DAYE: JO 982566:3
FILENG DAYE: JO 982566:3
FILENG DAYE: JOHN
NAME: KILKY JR. JOHN
NAME: KILKY JR. JOHN
NAME: ATTORNEY/AGENT INFORMATION:
NAME: KILKY JR. JOHN
NEGET JR. JOH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 1.5%; Score 18; Best Local Similarity 100.0%; Pred. No. Matches 18; Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 60601-6780
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPAIND:
OPERATING SYSTEM: PC-DOS/MS-L-L-S
SOFTMARE: Patentin Release #1.0, Version
APPLICATION NUMBER: US/05/091.332A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NERAL INFORMATION:
APPLICANT: KEENE, JACK D.
APPLICANT: KING, PETER H.
APPLICANT: KING, PETER H.
APPLICANT: LEVINE, TOOD
TITLE OF INVENTION: RECOGNITION, BINDING AND EXPRESSION OF RIBONUCLEIC ACIDS TITLE OF INVENTION: RECOGNITION, BINDING AND EXPRESSION OF RIBONUCLEIC ACIDS TITLE OF INVENTION: INVOLVED IN CELL GROWTH, NEOPLASIA AND INMUNOREGULATION NUMBER OF SEQUENCES: 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ent No. 5525495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION NUMBER: WO 97JP3438
APPLICATION NUMBER: WO 97JP3438
FILING DATE: 26-SEP-1997
435
ZIP: 22202
DMPUTER READABLE FORM
                                                                                                                                                  DDRESSEE: Овьин, ст.....
UDRESSEE: P.C.
TREET: 1755 Jefferson Davis Highway, Fourth Floor
                                                                                                                                                                                                                                                                                                     OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTALI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4; Length 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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ATTORNEY/AGENT INFORMATION:
AME: Oblon, No. 555495man F.
REGISTRATION UNMER: 24,618
REFERENCE/ACKET NUMEE: 714-156
TELECOMMUNICATION INFORMATION:
TELERONE: (703)13-3000
TELERA: (703)413-3000
TELERA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-TAUS
SOFTMANE: PatentIn Release #1.1, Version #1.25
REDII AFFLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JS/08/120,827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            714-156-0 CIP
```

Query Match 1.9%; Score 16; DB 1; Best Local Similarity 100.0%; Pred. No. 38; Matches 18; Conservative 0; Mismatches Length 57;

PPLICANT: KEENE, JACK D. PPLICANT: KING, PETER H. PPLICANT: LEVINE, TODD 5, Application US/08478675

ATPLICAMI: LEVINE, TODD

ITILE OF INVENTION: METHODS AND COMPOSITIONS USEFUL IN THE ITILE OF INVENTION: RECOGNITION, BINDING AND EXPRESSION OF ITILE OF INVENTION. INVOLVED IN CELL GROWTH, NEOPLASIA AND CORRESPONDENCE ADDRESS:
ACTURESSPONDENCE ADDRESS: RIBONUCLEIC ACIDS IMMUNOREGULATION

ALDRESSEE: P.C.
SIREE: 1755 Jefferson Davis Highway, Fourth Floor
CITY: Arlington
STATE: Virginia
COUNTRY: U.S.A.
21P: 22202

COMPUTEN ACCOUNTS.

COMPUTEN TYPE: Floppy disk
REDIN TYPE: Floppy disk
COMPUTEN: IBM FC compatible
COMPUTEN: IBM FC compatible
COMPUTEN: IBM FC COMPATIBLE
COMPUTEN: PC-DOS/MS-UGS
CORRENT APPLICATION COMPATIBLE
COMPUTEN: PC-DOS/MS-UGS
ETILING DATE: O7-JUN-1956
CILASSITICATION UNDER: US/08/478,675
FILING DATE: O7-JUN-1956
FRICAR APPLICATION LOATA:
APPLICATION UNDER: US 08/120.627
FILING DATE: 15-SEP-193
ATTORNEY/ACENT INDER: US 08/120.627
FILING DATE: 15-SEP-193
ATTORNEY/ACENT INDER: A18
REGISTANTION NUMBER: 40.18
REGISTANTION UNDER: 714-158-0 CIP
TELECOMMUTCATION UNFORMATION:
TELECOMMUTCATION UNFORMATION:
TELECOMMUTCATION UNFORMATION:

Page 10

HOLECULE TYPE: CDNA
US-08-478-675-85

TELEEX. (703)413-2220
TELEX: 248855 OPAT UR
FORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENUTH: 57 base pairs:
TYPE: nucleic acid
STRANDEDNESS: unknown
TOPOLOGY: unknown

85:

Ouery March 19%; Score 18; D6 1; Length 57; Best Local Similarity 100.0%; Pred Mo. 36; Marches 18; Conservative 0; Mismatches 0; Indels

```
NAME: BEOMO, SCOLT A.

REGISTRATION NUMBER: 32,724

TELECOMMUNICATION LAFORMATION:
TELEPHONE: (617),498-824
TELEPHONE: (617),846-881
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 308 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
Ouery March 1.9%; Score 18; DB 2; Length 308; Best Local Similarity 100.0%; Pred. No. 36; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: LAVAILES, Edward
APPLICANT: Recie, Lisa
APPLICANT: Recie, Lisa
APPLICANT: Recie, Javid
APPLICANT: Treaty, Javid
APPLICANT: Treaty, Hurice
APPLICANT: Spauldiny, Vikit
TITLE OF INVENTION: ENCODING THEM
NUMBER OF SEQUENCES: TROUBLES
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
STREET: 8D Cambridge
STATE: MASSachusetts
COUNTRY: Cambridge
STATE: MASSachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21F: 02140
COMPUTER REABLE FORM:
WEDLUH TYPE: Floppy disk
COMPUTER: HA PC Compactible
COMPUTER: HA PC COMPACTIBLE
COMPUTER: HA PC COMPACTIBLE
SOFTWARE: PATENTIA RELEASE #1.0, VETSIC.. #1.30
CURRENT APPLICATION DATA:
APPLICATION MURBER: US/08/721,488
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPLICANT: Jacobs, Kenneth
PPLICANT: MCCoy, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                    NEY/AGENT INFORMATION: E: Brown, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08721488
965388
                                                                                                                                                                                                                                               TELEPAX: 215-567-2991
TELEX: d011-48.
INFUMATION FOR SEQ ID NO: 157:
SEQUENCE CHARACTERISTICS:
LENGIH: 343 base pairs
TYPE: nucleic acid
STRANGENESS: double
TOPOLOT: linear
MOLECULE TYPE: DNA (genomic)
S-08-943-731-157
                                                                                                                     very Match 1.54; Score 18: Lb 4; Length 343;
Best Local Similarity 100.08; Pred, No. 35;
Matches 18: Conservative 0; Mismatches 0; Indels
```

0; Indels

0 Gaps

> 0 Gaps

```
STREET: FILL

STREET: FILL

STREET: PHILADELPHIA

COLYTY: PHILADELPHIA

STATE: PA

COUNTRY: USA

COUNTRY: USA

COUNTRY: USA

COUNTRY: USA

COUNTRY: PEE: PLOPPY disk

COUNTRY: PEE: PLOPPY disk

COMPUTER READABLE FORMPALLISH

COMPUTER READABLE FORMPALLISH

OPERATING SYSTEM: PC-DOS/MS DOS

SOTTMARE: PEE: PC-DOS/MS DOS

SOTTMARE: PC-DOS/MS DOS

SOTTMARE: PC-DOS/MS DOS

SOTTMARE: PC-DOS/MS DOS

SOTTMARE: PC-DOS/MS DOS

CURRINT APPLICATION NUMBER: US-06/212,322

APPLICATION NUMBER: US-06/212,322

APPLICATION NUMBER: US-08/212,322

APPLICATION NUMBER: US-08/212,322

FILING CATE: 14-MAR-1954

FILING CATE: 14-MAR-1954

FILING CATE: 03-DEC-1951

APPLICATION NUMBER: US-07/803,626

APPLICATION NUMBER: US-07/803,626

FILING CATE: 03-DEC-1951

INAME: DOTE LEARY Ph.D. KATHRYN

REDISTRATION NUMBER: 9508-27

TELEPHONE: 215-963-1284

TELEPHONE: 215-967-2991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LICANT: MARKOKO, LEBA, et al.

LE OF INVENTION: COMPOSITIONS AND METHODS FOR DETECTING

LE OF INVENTION: ALTEKED TYPE I OR TYPE IX COLLAGEN GENE SEQUENCES

BEROVEDING ALTEKED TYPE I OR TYPE IX COLLAGEN GENE SEQUENCES

BEROVEDING ADDRESS:
ORESSEE: PARTICH SCHMARZE JACOBS & NADEL, P.C.
IRRET: FLR:
ONE COMMERCE SQUARE, 2005 MARKET STREET, 22ND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /, Application US/08943731
6265157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SFOTILA. COMETTA D.
SEVILAS, CONSTANTINGS D.
SEREDA, LARISA
LARISA, CARISA W.
PACK, MICHAEL
COLICE, ALAIN
EARLY, JARES
KORKKO, JARNO
KORKO, JARNO
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09-342-653-

Application US/09342653

```
APPLICANT: Caboon, Sabecea E.
APPLICANT: Railski, Antoni
ITILE OF INVENTION (bromatin Associated Proteins
ITILE OF INVENTION (bromatin
```

RESULT 28 US-08-858-830-1/c

Ouery Match: 1.9%; Score 16; DB 1; Length 432; Bast Local Similarity 100.0%; Pred, No. 35; Matches 18; Conservative 0; Mismatches 0; Indels

0; Gaps

907 ggcaaaaaaaaaaaaaa 924 {!!!!!!!!!!!!!!!!!!!! 312 GGCAAAAAAAAAAAAAA 295

Application US/08858830

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Query Match 1.9%; Score 18; Best Local Similarity 100.0%; Pred. No. Matches 18; Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
STREET: 87 CambridgePark Drive
CITY: Cambridge
                                                                                                                                                                                                                                                                                                  REDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC:POS/MS:DOS
SOFTWARE: PSICENTIN Release $1.0, Version $1.30
RRENT APPLICATION DAIR:
RRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LICANT: Treacy, Maurice
LICANT: Spaudding, Yikki
LE OF INVENTION: SECRETED PROTEINS AND POLYMUCLEOTICES
LE OF INVENTION: ENCODING THEM
BER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RPPLICATION DATA:
CATION NUMBER: 08/702,080
           Massachusetts
: U.S.A.
2140
                                                                                                                                                                                                                 oplication US/08858834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TON NUMBER: 32,724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ON NUMBER: US/08/858,830 TE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kenneth
                                                                                                                                                                Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Edward
                                                                                                                                   Edward
```

IE: New York

Felfe & Lynch 5 Third Avenue T: Reinhard Buttner
T: Brigitte Kaluza
INVENTION: MELANOMA-INHIBIIING FRUTEIN
F SEQUENCES: 24

Application US/08578649 770366

TER READABLE FORM
TYPE: 3.5 inc

inch 1.44 Mb storage diskette

PC-DOS

DN NUMBER: DE P 43 24 247.2 [E: 20-July-1993 ENT INFORMATION:

MBER: US/08/578,649 29-July-1994

TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO:
SEQUENCE CHRACTERISTICS:
LENGTH: 581 base pairs
TYPE: nucleic acid
STRANDEDNESS: single

```
Dequence 58, Application US/0538592

Patent No. 666234

APELICANI INFORMATION:
APPLICANI ENDEGE, MILSON O., ET AL.
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS: II
FILE REFERENCE: CCIONA - 50X IS /05/385,582
CURRENT APPLICATION NUMBER: US/05/385,582
CURRENT FILLING DATE: 1999-08-30

EARLIER APPLICATION NUMBER: 05/07/17,393
EARLIER FILLING DATE: 1999-01-27
EARLIER FILLING DATE: 1999-01-27
EARLIER FILLING DATE: 1998-01-21
EARLIER FILLING DATE: 1998-01-21
EARLIER FILLING DATE: 1998-01-21
SOFTMARE: FILLING DATE: 1998-01-21
                                                                                                                                                                                                                                                                                               FEATURE:

NAME/KEY: misc_feature

LOCATION: (1):..(582)

OTHER INCORMATION: n - A.1,C or G
US-09-385-982-98
                                                                                                                                Overy Match 1.vs; Score 18; Di 4: Length 5s2;
Best Local Stailarity 100.04; Pred. No. 35;
Matches 18; Conservative 0; Hismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ULT 31
09-385-982-98/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
0; Gaps
                                                                                                                           0;
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: FELIDAE:
: KAMZ/KEY: misc_feature
: KUZAITON: (1) ...(621)
: COTHER INTLANATION: n * A.T.C or G
US-05-328-111-572
                                                                                                                                                                                                                                                                                        APPLICANI: Derti, Adnan
APPLICANI: Ford, Donna M.
APPLICANI: Ford, Donna M.
APPLICANI: Mornham: John E.
APPLICANI: Mornham: John E.
APPLICANI: Schiegel, Kobert
ITILE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
ITILE OF INVENTION: PRODUCTS
FILE REFERENCE: CCD-257 (US)
CUBARNI APPLICATION UNUMERE: US/05/328.111
CUBARNI FELICATION UNUMERE: US 60/088.801
EARLIER APPLICATION UNUMERE: US 60/088.801
EARLIER APPLICATION UNUMERE: US 60/088.801
                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 850
SCITWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 572
                                                                                                                                  CRIENISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09326111
                                                                                                                                                                                                                                                                                בבת: US 60/088,601
98-06-10
```

NAME/KEY: mat\_peptide; ; LOCATION: 179..459 US-08-578-649-4

NAME/KEY: sig\_peptide LOCATION: 110..178 E/KEY: CDS ATION: 110..499 LOGY: linear LE TYPE: cDNA

Overy Match 1.9% score 18; Lb 1; Length 581; Best Local Similarity 100.0%; Pred. No. 35; Matches 18; Conservative U: Hismatches O; Indels

```
APPLICANT: Allen, Steve
APPLICANT: Rafalski, Antoni
APPLICANT: Sakai, Hajime
TITLE OF INVENTION: Plant Hetal Transporters
FILE REFERENCE: BBJ303 US WA
CUBARNI APPLICATION NUMBER: US/05/461,474
CUBARNI FILING DATE: 1999-12-14
DAKLIER APPLICATION NUMBER: 60/112,562
EAKLIEK FILING DATE: 1998-12-16
: IYFE: DNA
: OKJAWISM: Triticum mestivum
US-09-461-474-13
                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Microsoft Office 97
SEC ID NO 13
LENGTH 790
TYPE: TANA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 1.9%; Score 18; Best Local Similarity 100.0%; Pred. No. Matches 18; Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                  Application US/09461474
78042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sdps
```

Length 621;

Query Match 194; Score Hs; DB 4; Length 790; Best Local Similarity 100.0%; Pred. No. 34; Matches 1B; Conservative 0; Mismatches 0; Indels

Gaps

5, Application US/08684672 5700926

```
APPLICANT: SCHLESSINNER, David
APPLICANT: de la CHAPELLE, Albert
APPLICANT: SRIVASTAVA, ANDAUG KUMBA'
TITLE OF INVENTION: MOLECULAR CLOUING OF THE ANHIUMOI
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE, SHECKER & MAIHIS, L.L.P.
STREET: P.O. BOX 1404
CITY: Alexandria
STATE: Viginia
COUNTRY: Unital
                                                                                                                                                                                                                                                                                                                                                                                                                         ; NAME/KEY: CDS
; LOCATION: 243..650
US-08-684-672-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC POS/MS-DOS
OPERATE: PATENTION RELEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U5/08/664.672
FILING DATE: 22-UL-1956
CLASSIFICATION NUMBER: U5 08/052,937
PRICH APPLICATION NUMBER: U5 08/052,937
FILING DATE: 27-APR-193
ATTORREY/ACENT INFORMATION:
APPLICATION NUMBER: U5 08/052,937
FILING DATE: 27-APR-193
ATTORREY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                           Query Match 1.9%; So
Best Local Similarity 100.0%; I
Matches 18; Conservative 0;
                                                                                                                                                                                                                        TELEFAX: (703) 836-202
FORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 854 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: O'Shaughnessy, Brian P.
REGISTRATION NUMBER: 32,747
REFERENCE/DOCKET NUMBER: 0305-
PLECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
PPLICANT: KERE, Juha
PPLICANT: SCHLESSINGER, David
PPLICANT: de la CHAPELLE, Albert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22313-1404 States
                                                                                                             Application US/08684672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /A. Anand Kumar
MOLECULAR CLONING OF THE ANHIDMOTIC
ECTODERMAL DYSPLASIA GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     030956-002
                                                                                                                                                                                                                                                                                                                                      DB 1:
                                                                                                                                                                                                                                                                                                           Gaps
```

Application US/08896410

FEAL INFORMATION:
PFILCANI: The Government of the United States of America.
PFILCANI: as represented by the Secretary, Department of Health and Human FFILCANI: Services
TILE OF INVENTION: Mammalian Muscle NAD:Arginine
TILE OF INVENTION: ADP-THOSYLtransferase
NEER OF STORMENCES: 36

NDDRESSEE: SIREET: 62 CITY: Newp STATE: Cal

ifornia

E: Knobbe, Martens, Ölson & Bear 620 Newport Center Drive, l6th floor ∺Mport

ESPONDENCE ADDRESS:

FUTER KRADABLE FORM:
EDIUM TYPE: FLOPY disk
CARUTER: IBM PC compatible
PERATURS SYSTEM: PC-DOS/MS-LOS
VOTIMARE: PRIEMENT RELEASE #1.0, Version #1.25
MEENT APPLICATION DATA:

```
; HULECULE TYPE: DNA (genomic)
US-08-644-672-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ALCRESSEE: BURNS, DOANE, SWECKER & MATHIS, L.L.P.
SIRET: P.O. BOX 1404
CITY: A lexandria
STATE: Virginia
CUNNINY: United States
2 LP: 22313-1404
CLAPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CCMFUTER: IBM FC COMPAILDIC
CFEATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0. Version #1.30
CUREENI APPLICATION MUMBER: US/08/684,672
FILING DATE: 22-JUL-1956
CLASSIFICATION: 435
Oder; Match 1.9%; Score 16; DB Best local Similarity 100.0%; Pred. No. 34 Matches 18; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/052,957
FILIG DATE: 27 AFF-1593
ATJORNEY/AGENT INFOMMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AFFILEANI: SRIVASIAVA, AMEND KOMET

IILE OF INVESTION: MALECULAK CLOSHNG UF THE ANHIDROTIC

RIVER OF INVESTION: ECIODERMAL DYSPLASIA GENE

KIMER OF SECUENCES:

CAMERSFORIZMOE ALDRESS:
                                                                                                                                                                                                                                                                                                                                                             A.AME: O'Shaughnessy, Brian REGISIRATION NUMBER: 32,747 REFERENCE/DOCKET NUMBER: 03 RECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                    030956-002
                                             Length 854;
   Gaps
```

APPLICATION NUMBER: US/08/896,410 FILING DATE: 18-JUL-1997 CLASSIFICATION: 435

```
NAME: ANILO M. KIFSAITICK
REGISTRATION NUMBER: 321617
REFERENCE/DOCKET NUMBER: NIH039.001DVI
TELECHOME: 619-33-3550
TELEPHONE: 619-33-3550
INCOMMATION FOR SED ID NO: 37:
SERVING: CHANCETER SITES:
SERVING: COS
MARCET COS
MARCET
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; FEATURE:
; NAME/KEY: CDS
; LOCATION: 1..855
US-08-631-200-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; NAME/KEY: CDS
; LOCATION: 2..1039
US-05-073-257-1
                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-UOS
SOFTWAKE: PATENTIAN RELEASE #1.0, VETSION #1.30
CURRENT APPLICATION DATA:
APPLICATION UNBER: US/08/631,200
FILING DATE: 12-APR.1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
ANAPT: COTIFY! INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 1.9%; Score 16; Best Local Similarity 100.0%; Pred. No. Matches 16; Conservative 0; Mismatc
                                                                                                                                                                                                                             NAME: COTUZZI, LAUTA A.
REDISTATION NUMBER: 30,742
REPERRNEZ/POCKET NUMBER: 7853-
TELECOMUNICATION: TORBATION:
TELEFANE: (212) 790-900
TELEFANE: (212) 69-9741/8864
TELEFAN: 6141 FENIE
                                                                                               TYPE: fucleic acid
STRANTEINESS: unv
                                                                                 TOPCLOGY: unknown
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pence 14, Application US/08631200 ent No. 5646040 NERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PLICANI: Kleyn, Patrick W.
PLICANI: H.Ore, Karen J.
TIE OF INVENTION: COMPOSITIONS FOR THE IREATMENT AND
TIE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TIE OF INVENTION: DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY
MBER OF SEQUENCES: 59
WRESPONDERCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGIA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NSTH: 1333 base pairs
PE: nucleic acid
RANDEDNESS: single
                                                                                                                                                                                                                                                                                                                 7853-057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1333;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
```

Query March. 1.9%; So Best Local Similarity 100.0%; I Matches 18; Conservative 0;

0; Indels 0;

Gaps

Length 1338;

907 ggcaaaaaaaaaaaa 924

```
; NAME/KEY: CDS
; LOCATION: 1..855
US-08-829-553-14
                                                                                                                                                                                                                                                                                                                 usery Match 194; score 18; DB 1; Lengtl 1338; Best Local Similarity 100.0%; Pred. No. 33; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: COTUZZI, LANTA A.

REGISTRATION NUMBER: 30.742
REFERENCE/DOCKET NUMBER: 7653-0
REFERENCE/DOCKET NUMBER: 7653-0
RELEPHONE: (212) 790-9590
TELEPAX: (212) 869-9741/8864
TELEXA: 6641 PENNIC
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 1318 base pairs
TYPE: NUCLEIC CAI
STRANDEDNESS: UNKNOWN
STRANDEDNESS: UNKNOWN
                                                                                                                                                                                                                                     STREET: 1155 Avenue of the Americas CITY: New York
STRTE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
MAPUTEP 10036-2711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TIPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/825,553
EILING DATE: J8-MAR-1997
CLASSIFICATION: 530
PRICE APPLICATION DATA: 500
APPLICATION NUMBER: US 08/631,200
FILING DATE: IZ-APP-1996
ATTORREY/AGENT INFORMA: ION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LICANT: MODTE, KACHI JEMEN AND LE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND LE DE INVENTION: DIAGNOSIS OF BODY WEIGHT DISORDERS. 59
RESPONDENCES: 59
RESPONDENCE AUDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/08829553
                                                                                                                                 Application US/08922267A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unknown
E: DNA (genomic)
PENTION: COMPOSITIONS FOR THE TREATMENT AND TEMPOSITY OF BODY WEIGHT DISORDERS, INCLUDING OBESITY NUMBERCES: 82
                                                      leyn, Patrick W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26 MAR-1997
N: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7853-057
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cery match 1.9%; Score 18; DB 2; Length 1338; Best Local Similarity 100.0%; Pred No. 31; Matches 18; Conservative 0; Mismatches 0; Indels

Gaps

0.

LI 41 6-936-707A-14 quence 14, App

1, Application US/08936707A 5871931

INFORMATION:
CANT: Kleyn, Falrick W.

CANT: Kleyn, Falrick W.

CANT: MOOTE, Karen J.

COMPOSITIONS FOR THE TREATMENT AND

OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND

OF INVENTION: DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY

R OF SEQUENCES: 60

: Pennie & Edmonds LLP 1155 Avenue of the Americas

IBM PC compatible

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SIREPI: 1155 AVENUE OF the Americas
CIIY. New York
COUNTRY: U. 5.A.
COMPUTER READABLE FORM:
MEDIDM TIPE: Floopy disk
MEDIDM TIPE: Floopy
MEDIDM TIPE: MARE: GOID TO NO. 14:
SECURACE CHARACTERISTICS:
LENGTH: 1336 BASE PAIRS
SECURACE CHARACTERISTICS:
LENGTH: 1336 BASE PAIRS
MARE/KN: CDS
MARE/KN: CDS
MARE/KN: CDS
MARE/KN: CDS
MARE/KN: CDS
MARE/KN: CDS
US-GB-92-2670-14
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OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30

ION DATA: MBER: US/08/936,707A 24-SEP-1997

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; NAME/KEY: CDS
; LOCATION: 1..855
US-08-936-707A-14
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Best Local Similarity 100.0%;
Matches 18; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NTTORNEY/AGENT INFORMATION:
NAME: COLUZZI, Laura A.
REGISTRATION NUMBER: 30,712
REFERENCE/DOCKET NUMBER: 7853-
TELECOMMUNICATION INFORMATION:
TELECHONE: (212) 790-9990
TELEFEX: (211) 869-9741/8864
TELECEX: (211) 196NUE
                                                                                                                                       RENT APPLICATION DATA:
PLICATION NUMBER: US
LING DATE: 24-SEP-19
                                                                                                                                                                                  DIUM TYPE: Floppy disk

MPUTER: IBM PC COMPATIBLE
ERATING SYSTEM: PC:DOS/MS-DOS
FTWARE: Patentia Release #1.0, Version #1.30
FWT ARPHIPATYN DAYA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANDEDNESS: of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08936706A
5876919
                                                                                                                                                                                                                                                                                                                                       Pennie & Edmonds LLP
                                                                                                                 24-SEP-1997
N: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER: 30,742
CKET NUMBER: 7853-100
                                                                                                                                                                                                                                                                                                                                                                                        N: COMPOSITIONS FOR THE TREATMENT AND N: DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY 35: 60
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                                                     853-055
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: NAME/KEY: CLS
: LOCATION: 1..855
US-D8-936-706A-14
: LOCATION:
US-09-248-203-14
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                                                                                                    TELEFYAX: (ZIZ) 865-9741/8864
TELEX: 66141 PENNIE
SEQUENCE CHARACTERISTICS:
LEKKTH: 1336 base pairs
TYPE: INCLEASE acid
STANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Pennie 6
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: 1EM PC compatible

OPERATING SYSTEM: PC-DOS/MS-LOS

SOPTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Rel
                                                                                                                                                                                                                                        REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 78
LECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                          PLASSIFICATION:
IOR APPLICATION DATA:
US 08/936,707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     THE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND THE OF INVENTION: DIAGNOSIS OF BODY WEIGHT DISORDERS, BER OF SEQUENCES: 60
                                                                                                                                                                                                                                                                                                          ILING DATE: 24-SEP-1997
ORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               ICATION NUMBER: US/05/248,203
NG DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
DMESS: single
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                                                                                                  single
                                                                                                                                                                                                      969-9741/8864
                                                                                                                                                                                                                                                           7853-160
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.very Match . 1.5%; Score 16; Db 3; Length 1338; Best Local Similarity 100.0%; Pred No. 33; Matches 16: Conservative 0; Mismatches 0; Indels

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; NAME/KEY: CDS
; LOCATION: 1..855
US-09-406-071-14
                                                                                                                                                                                                                      Ouery Match 1.94; Score 18; DB 4; Length 13:c; Best Local Similarity 100.04; Pred: No. 33; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING JATE:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, Laura A.
REGISTRATION NUMBER: 30,742
REGISTRATION NUMBER: 7853-1
TELEPOMORIFICATION INFORMATION:
TELEPOMORIFIC (212) 790-9909
TELEPAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 66141 PENNIE

NFORMATION FOR SEQ ID NO: 14:

SEQUENCE CHARACTERISTICS:

LENGTH: 1338 base pairs

TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACIBLE
OPERATING SYSTEM: PC COS/MS COS
SOFTWARE: PATCHILD Release 1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPLICANT: MOORE, KAREN J.
WITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
WITLE OF INVENTION: DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY
WHORE OF SEQUENCES: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4, Application US/09406071 6207386
PORMATION:
T: Kleyn, Patrick, and Moore, Karen
TINVENTION: RP Compositions and Therapeutic and
INVENTION: Diagnostic Uses Therefor
F SEQUENCES: 16
                                                                                                        Application US/08955918C
168130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pennie & Edmonds LLP
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ON NUMBER: US/09/406,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
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                                                                                                                                                                                                                                                                                               0; Indels 0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Search completed: April 30, 2002, 10:41:30 Job time: 10155 sec
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; NAME/KEY: CDS
; LOCATION: 1..858
US-08-955-918C-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:

CORRESPONDENCE ADDRESS:

ADDRESSEE: LAHIVE & COCKFIELD, LLP
SIRET: 28 State Street

CITY: BOSTON

STATE: BASSACINSELLS

COUNTRY: USA

LIP: 02109-1875

COMPUTER READABLE FORM:

COMPUTER: LBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: LBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: LBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: LBM PC compatible

OPERATION SYSTEM: PC-DOS/MS-DOS

SOFTMARE: LBM PC compatible

OPERATION SYSTEM: PC-DOS/MS-DOS

SOFTMARE: LBM PC-COMPATION

APPLICATION UNMERE: US/OB/955,518C

FILING DATE: 29-MG-1996

ATTORREY/ACENT INFORMATION:

NAME: MADDRESSEE: US/OB/957,766

ATTORREY/ACENT INFORMATION:

NAME: MADDRESSEE: MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/MS-DOS/
                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
TELEFAX: (617)227-5941
TORRAKTION FOR SEO ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 1338 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Mandragouras, Amy E.
RECISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MNI
LECONGUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MNI-007CPDV2CPA
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Overy Match 194; Score 18; DB 4; Length 1338; Best Local Similarity 100.0%, Pred: No. 3; Matches 18; Conservative 0; Mismatches 0; Indels 0; Gaps

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Page 19

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Title:
Perfect score:
Sequence:
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Run on:
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R_Geneseq_1101:

| Continue | Receiver | Rec
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Copyright (c) 1993 · 2000 Compugen Ltd.
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Homan secreted pro	AAV34234	5	500	3./	2.2	こうべき
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com achacuce or t	116767	د	7331	77 0	7.0	0
CDNA sequence of h	AAC79442	2	590	100.0	0.60	
DNA encoding human	222000				9 0	, ,
	LICTBARK.	30	560	100.0	590	-
Description	ID	DB	Match Length DB	Match	Score	No.
				Query		Result

PHX

New breast tunour protein genes used, in vaccines for immunotherapy, or for diagnosis of breast cancer

Human lung tumour	AA208301	20	278	₩ •	20
Plant microsatell	AAA31526	21	242		20
	AAA61347	21	115	4	20
Human novel secre	AAV26706	19	79	-	20
Model mRNA #3 used	AAA61354	21	51		20
Human mitochondria	AAT17027	17	4.2	4	2
	AAX76360	20	40		20
Pneumocystis carin	AAX76366	20	U.		200
Human cancer assoc	AAC77811	21	5830		1
Sequence encoding	AAN50530	0	5/60		1.2
Renal cancer assoc	AAX35681	20	5306		2.
S.	AAH19355	22	1685		21
Eph-related tyros	AAQ90652	16	3133		21
Human breast tumou	AA233647	20	2281		21
Honeybee alpha-glu	AAH48294	22	1994		22
polynuc	AA158747	22	1763		21
sativa pi	AAQ85984	16	1624	3.6	21
secreted	AAD08465	22	1549		21
ortholog	AAC91065	22	1547		21
secreted	AAD11689	22	1373		21
secreted	AAZ98076	21	1373		21
cDNA encoding memb	AAA07076	21	1302		21
Nucelotide sequenc	AAV53733	19	1302		21
DNA encoding endot	AAT17380	17	1302		21
colon c	AAH34582	22	1279		21
cancer as	AAC78075	21	1279		21
	AAC 934 27 ·	21	1164		21
secreted	AAX20466	20	1070		21
Human secreted pro	AAF22420	21	1649		21
otide sequ	AAV59132	19	959		21
transfer pr	AAG74410	15	679		21
Probe #2 used in a	AAF50545	22	67		21
Human kinesin-like	AAC66548	21	121162		22
Human neuroblastom	785	22	3448	3.7	2

AAX35  AAX66  AAX66  On Si  DNA e  DNA e  Frees  Brees  Bress  Cress  10-7-11  17-11				×
AAX64213 standard; CDNA: 590 BP.  AXX64213;  Object 1999 (first entry)  DNA encoding human breast tumour protein imm Breast tumour protein; immunogenic fragment; Creast cancer development; therapy; ss.  incospiens.  Krassid66-A2.  verjUL-1996; 96NO-USZ7416.  17-UUL-1996; 96NO-USZ7416.  17-UUL-1997; 97US-0948255.  24-EEC-1997; 97US-0948255.  17-UL-1996; 96US-0118594.  (CCRI-) CORIAA CORP.			Reed SG, Xu J	P X
AAAX64213 standard; CDNA; 590 BP.  AAX64213;  Dr5EP-1999 (first entry)  DXA encoding human breast tumour protein imm  Breast tumour protein; immunogenic fragment;  breast cancer development; therapy; ss.  ELLS sapiens.  KLS533665-A2.  US-JUL-1996; 96WO-US27416.  17-JUL-1996; 96WO-US27416.  17-JUL-1997; 97US-0998253.  24-DEC-1997; 97US-0998253.  24-DEC-1997; 97US-0998253.  21-DEC-1997; 97US-0998253.  21-DEC-1997; 97US-0998253.  21-DEC-1998; 96US-0118554.		CORP.	(CURI-) CORIXA	P
AAX84213 standard; CDNA; 590 BP.  AAX84213;  Un-SEP-1999 (first entry)  DNA encoding human breast tumour protein imm  Breast tumour protein; immunogenic fragment;  breast cancer development; therapy; ss.  number sapiens.  Augustus Sapiens.				X
X4213 X4213 XAX6213; AXX6213; On-SEP-1999 (first entry) DNA encoding human breast tumour protein imm Breast tumour protein; immunogenic fragment; Ereast cancer development; therapy; ss. HLDS aspiens. HLDS 33865-A2. Un-JUL-1996: 98WO-US27416. 17-JUL-1996: 96WO-US27416. 17-JUL-1996: 96WO-US27416. 17-JUL-1996: 96WO-US27416. 17-JUL-1996: 96WO-US27416.	554.	9805-0118		אל
AAX84213 standard; CDNA; 590 BP.  AAX84213;  Un-SEP-1999 (first entry)  DNA encoding human breast tumour protein imm  Bréast tumour protein; immunogenic fragment; breast cancer development; therapy; ss.  human sapiens.  AUX9533869-A2.  Un-301-1999; 21-Lic-1998; 98W0-US27416.  17-JUL-1998; 98W0-US27416.  17-JUL-1998; 98W8-US27416.	255.	970S-0998		תו או
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Least Cancer; Vaccine; ss.
The present sequence was isolated from a presst tumour cDNA library. It a provided in a specification relating to compounds for immunotherapy and diagnosis of breast cancer. Breast tumour antigens and the injunction of the used in the preduction of the used in the treatment of breast cancer. Soliferated composition to be used in the treatment of breast cancer. Soliferated peells and incubated antigen presenting cells are also equivalence a vaccine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention provides the protein and coding sequences of a number of colorectal cancer antiques. These are shown in the AA157547-AA157549 and AA438565-AA436641. These can be used in the diagnosis, prevention and treatment of cancer of the colon and/or rectum. The present sequence is a colorectal cancer antiqen general sequence. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from HIFO at fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated polypeptide for treating, preventing and/ or progressing disorders related to the colon and rectum including colorectal cancers and also for testing and detection e.g. diagnosis
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Human, colorectal cancer; colorectal cancer antigen; gene therapy; ss. Auman colorectal cancer antigen cDNA SEQ ID NO: 59.

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2000US -0212467

2000US -021467

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17 - KOV - Zuvu
18 - CEC - Zuv
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                                                                                                                                                                                                                                                                                                                               Claim 1; SEv ID NO: 59; 522pp - Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI: 2001-457727/45.
P-PSDB: AAM38617.
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The present invention provides the protein and coding sequences of a number of colorectal cancer antigens. These are shown in AAISTS47-AAISTS41. These can be used in the disquests, prevention and treatment of cancer of the colon and/or rectum. The present sequence is a colorectal cancer antigen coding sequence of the invention.

The invention are the colorectal cancer antigen coding sequence of the invention.

The sequence data for this patent did not term part of the printed specification, but was obtained in electronic format directly from WIFO at fig.-VIPO-int/pub/published\_pot\_sequences. Sequence 629 BP; 244 A; 94 C; 79 G; 212 T; 0 other;

Query Match Best Local S Matches 427 10.0%; Solution 72.4%; Solution 72.4%; Solution 100.0%; F 427; Conservative 6; Score 427; DB 22; Length 6; Pred. NO: 1.6e-150; 0; Mismatches 0; Indels Length 629; 0; Gaps

2 5 2 8 E) 121 LUCCCCUJ95LLAGCAGITTAAATGAAACAGAGITCATCAATGAAATGAGATATITAAAA 259 cl trabajecettigigecetgiticoettigigecetabititicoparaet 120 37a TIAAAGSCIGIIGIGACCAIGITICCCITIGIAGCAAIAAAIGITITITIACGAAAACT 319

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Matches 23
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21-DEC-1999;
14-MAR-2000;
12-MAY-2000;
09-JUN-2000;
21-JUL-2000;
                                                                                                                                                                                                          The invention relates to movel genes (Ahhö)77-Ahfijii) associated with cervical cancer with cytostatic activity. The nucleic acids and encoded polypeptides are useful: to assess if a patient is affilicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for inhibiting cervical cancer in a patient. The nucleic acids may also be useful for gene thorapy.
                                                                                                                                                                                 Sequence 468 BP; 121 A; 80 C; 95 G; 168 T; 4 other;
                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 120-121; 1051pp: English.
                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid for diagnosing and treating cervical cancer and for assessing and detecting compounds for treating the cancer .
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                                                     08-DEC-2000; 2000WO-US33312
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78 TIGTTCCTTTTCCCTGTGCCTGTGTCAAATCTTCAAGTCTTCAAGTCTGAAAATACATTTGATA 19
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                                                                                                           h 3.9%; So Similarity 100.0%; p 23; Conservative 0:
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990S-0171350.
2000US-0189315.
2000US-0203791.
2000US-0210600.
2000US-0220114.
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Pred. No. 6.6:
Mismatches U: Indels
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RESULT 6 AAH70254/c

Sytuatatic; immunosuppressive; nourcepts; neuroprotectitye; antiviral; antiallergic; hepatotropic; antidiaketic; antificilammatory; antiviral; vulidrary; anticonvulsant; antibacterial; antificingal; antiparasitic; cardiunt; gene therapy; cancer; immune disorder; cardiovascular disorder; cardiovascular disorder; cardiovascular disorder; secreted protein; ss.

Human secreted protein cDNA #10.

22-HEB-2001 (first entry)

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RESULT 7
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ID AAC$6505 standard; CDNA; 638
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Matches 23
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12-MAY-2000;
05-JUN-2000;
21-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to novel genes (AMH6877-AMH7383) associated with cervical cancer with cytostatic activity. The nucleic acids and encoded polypeptides are useful: to assess if a patient is afflicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a pre-malignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for industring cervical cancer in a patient. The nucleic acids may also be useful for gene therapy.
                                                                                                                                                                                                                                                                                                                                            Sequence 543 BP; 134 A; 103 C; 117 G; 189 T; 0 other;
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21-2EC-1999;
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23: Conservative
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990S-0171350.
20060S-0189315.
20060S-0203791.
20060S-0210600.
20060S-0220114.
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Pred. No.
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48 human secreted proteins AABS2104-BS2180. The genes can be used to generate fusion proteins AABS2104-BS2180. The genes can be used to generate fusion protein so y linking to the gene for the human (Immunoylobulin G FC portlon (SEQID) for increasing the stability of the fusion protein as compared to the human protein only. The genes and proteins are useful for preventing, amaliorating or treating medical conditions, e.g. by protein or gene thrappy. The genes are isolated from a range of human tissues disclosed in the specification. The diagnosis treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer, and other cancers of the address gland, bone, bone and ovarian cancer, and other cancers of the address gland, bone, bone amonytic anaems, autoimment thyroiditis, diabetes mellitus, Crohn seconds, autoimment thyroiditis, diabetes mellitus, Crohn seconds, (c) cardiovascular disorders such as mycoardial ischaemias; (d) cond healing; (e) neurological diseases e.g. cerebral aroxia and condition infections.
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Best Local S
Matches 22
Human: secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative discrete; pleukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; inflammate; system; asthma; lymphocytic disease; brain; heratic; lymphocytic disease; brain; heratic; lymphocytic, fundamation; ischaemic shock; Alzheimer's disease; restecosis; AIDS; cognitive disorder; schizophrenia; prostate; obesity; ustooclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;
                                                                                                                                                                                                                                                                                      Human secreted protein gene 16 clone HMABH07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV34234;
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20-JAN-2000; 2000US-0176930.
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P-PSDB; AAB52112.
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В οy

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Matches 22
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Feng P, F
Kyaw H, L
Ruben SM,
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22-AUG
        New isolated human genes and the secreted polypeptides they encode useful for disquosis and treatment of e.g. concers. it. closical disorders, immune diseases, inflammation or blood disorders
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P-PSDB; AAW75137.
                                                                                                                                             sh 3.7%; so smilarity 100.0%; F 22; Conservative 0;
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                                                                                                                                             Score 22; DB; Pred. No. 14; 0; Mismatches
                                                                                                                                                                                DB 15;
14;
                                                                                                                                                                                                                                                                                                                209 T; 3
                                                                                                                                                 <u>.</u>.
                                                                                                                                                                                                               Length 1.53;
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Hu JS;
Rosen CA;
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AVVISUAL 5

AVVISUAS Standard; DNA; 1075 EP.

ACC AVVISUAS;

ACC ACCOUNTS SECRED PROTEIN; fusion provided for the following following for the following fol
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Best Local S
Matches 22
                11-SEP-1998
 06-MAR-1998;
                               MO9839445-A2
                                             Homo sapiens.
                                                                                                                                            28-JAN-1999
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                                                                                                                                                                                                                                                                             Sequence 1075 BP: 251'A: 308
                                                                                                                                                                                                                                      th 3.7%; Sc
1 Similarity 100.0%; p
22: Conservative 0;
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98WO-US04492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human: secreted protein: fusion protein: gene therapy; protein therapy; diagnosis: tissue: cancer; tumour: neurodegenerative disorder; leukaemia; develor-mental abnormality; foetal deficiency; blood; allergy; renal ds; inflar-ation; schemn; lymphocytic disease: brain hepatic; lymphoma; inflar-ation; schemn; Albeimgr disease; brain; schemn; Albeimgr; disease; restenosis; Albe; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteocrorsis; arthritis; testis; lung; thyrodditis; thyroid; disestion; endocrina; metabolism; regulation; malabsorption; gastritis; neoplasm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acid sequences: AAV34154-V34276: amino acid sequences AAW75057-W75179) which are useful for preventine, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polyproptides in a sample or by determining the presence of mutations in the new polyprolectides. Specific uses are described for each of the 70 polyprolectides, based on which tissues they are most highly expressed (see AAV34154 for described uses).
7 MAR 1997
1 APR 1997
protein dene 16 clone HMABH07
Score 22: DB 1: Pred. No. 14: 0: Mismatches
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This sequence represents a nucleic acid molecule which encodes a secreted human protein. The gene number, and the clone it is derived from, are detailed in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene to a human immanoglobulific portion (e.g. ANVA145) for increasing the stability of the fused protein as compared to the human protein only.

Section of the second

New isolated human genes and the secreted polypeptides they encode - useful for diagnosts and treatment of e.g. cencers, neurological disorders, immune diseases, inflammation or blood disorders

Beddarik DP, Brewer LA, Carter KC, Duan R, Ebner R, Feng P, Ferrie AM, Fischer CL, Graves KA, Greene JM, Kyaw H, Lafleur DW, Li Y, Moore PA, Ni J, Olsen HS, Ruben SM, Shi Y, Soppet DR, Young PE, Yu GL, Zeng Z;

Endress GA:
Hu JS;
Rosen CA:

( HUMA - )

HUMAN

GENOME SCI INC

Claim 1; Page 234-235; 447pp; English.

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This sequence represents a nucleic acid molecule which encodes a screteed himan protein. The game number, and the clone it is derived from, are detailed in the descriptor line. The game can be used to generate fusion proteins by linking to the game to human immunoglobulin CC Fc portion (e.g. ANV3145) for increasing the stability of the fused CP protein as compared to the human protein only.

CC The invention relates to 70 novel genes and their fragments (nucleic acid seguences: ANV31454 vy14276; amino acid sequences ANW5057-475179) which are useful for preventing, treating or ameliorating medical CC conditions e.g. by protein or gene therapy. Also, pathological CC conditions can be diagnosed by determining the amount of the new polynucleotides. Specific uses are described for each of the 70 CC the new polynucleotides. Specific uses are described for each of the 70 CC polynucleotides. Specific uses are described for each of the 70 CC polynucleotides based on which tissues they are most highly expressed in CC (see ANV31154 for described uses).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated human genes and the secreted polyreptides they encode useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
                                                             polyA_signal
                                                                                                                             misc_feature
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EP192392-A
                                                                                                                                                                                                                                                                                                     sig_peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence of human insulin receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-JUL-1991 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAN50009 standard: cDNA: 5198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 175; 447pp; English.
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P-PSDB: AAW75072.
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Puben SM, Shi Y, Soppet DR. Young PE, Yu GL, Zeng Z;
                                      /*tag- c /roduct beta subunit 278.2289 /*tag- d /roduct precursor processing site* 5160.5165 /*tag- e /*tag-
                                                                                                                                                                                           /*tag= b
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2290..4150
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                                                                                                                                                                                                                                                                                                  31-AUG-1999; 99JP-0245962
09-MAY-2000; 2000JP-0136266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; chromosome 1; 1935; neuroblastoma cell line: NB·1: anticancer; tumour suppressor; human 1p36 homozygosity delation domain; tumour; diagnosis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A mutant IR is claimed which may have a mutated alpha-chain, esp. at the precursor processing site. The beta-chain may be mutated, e.g. by deletion of the transmembram sequence; the tyrosine kinase activity may be inactivated. Fig. 5 is a comparison of oncogene and human EGF receptor sequences with that of HIR in the cytoplasmic domain of the insulin receptor beta subunit.
                                                                                                                 WPI; 2001-226686/23.
                                                                                                                                                                                                                                                                                                                                                                                   31-AUG-2000: 2000WO-JP05930
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                                                                                                                                                                                                                       (HISM ) HISAMITSU PHARM CO LTD. (CHIB-) CHIBA PREFECTURE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human neuroblastoma cell line NB-1 lp36 nucleotide sequence SEO ID NO:59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bell JR,
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diagnostic
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes the coding and protein sequences of the human kinesin-like protein iRLP. It is thought that the protein could be involved in neurological disorders, infortility, spontaneous abortion, neonatal chromosome disorders, aneuploidy and cancers. This is due to its function in the movement of microtubbles. The protein shows homology to the murine KIFIA and KIFIB proteins. The sequences disclosed in the invention can be used in the isolation of similar human proteins and in vector production. In addition, the biallelic markers shown can be used in disease diagnosis and population studies.
                                                                                                                                                                                                                             Claim 1: Page 143-175: 199pp: English.
                                                                                                                                                                                                                                                                 An isolated or purified human kinesin-like protein (HKLP) encoding polynucleotide in a sample comprises a contiguous span of at least 12 nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human: kincsin-like protein: HKLP: KIF1: cell division: cancer: intracellular transport: neurological disorder; infertility; biallelle marker: spontaneous abortion: neonatal chromosome disorder: aneuploidy: ds.
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                                                                                                                                                                                                                                                                                                                                                                                                           Bouqueleret L. Dufaure-Gare 1, Grel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a homogygosity deletion domain co-existing in the 35-position of the first chromosome short arm (1p35) in human neuroblastoma. Also described are base sequences from the 1p36 position of human neuroblastoma. Also described are base sequences (RR-1 and MASS-NB-SCH-1), which are tumour suppressor genes in human neuroblastoma. The genes are tumour suppressor genes, base sequence data of which are applicable as fumour markers and respents in studying mechanism of tumour body (ormation, and gene diagnosts of tumours as well as in developing anti-cancer drugs. AMS78787 to AMS78820 propresent PCR primers used in the exemplification of the present invention, and AMS7830 to AMS78821 prepresent sequences given in the exemplification of the present invention of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-OCT-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-FEB-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ch 3.7%; Score 22; DB 1 Similarity 100.0%; Fred. No. 8.1 22; Conservative 0; Mismatches
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Sequence 121162 BP; 33272 A; 24108 C; 25842 G; 37919 T; 21 other:

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RESULT 14
AAF60545
RESULT 15
AAQ74410
ID AAQ744
XX
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Best Local Similarity 100.0%; Pred, No. 7.5;
Matches 22; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                    Ouery Match 3 1.61, Score 21; DB 22; Length 57
Best Local Similarity 100.01, Pred, Mo. 48;
Matches 21: Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ74410 standard; DNA; 679 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting and quantitating analyte polynucleotide in a sample, by co-amplifying analyte polynucleotide with predetermined arount of pseudo target, producing amplification products and quantifying analyte amplicons
                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a method for quantifying analyte polynucleotides (AP). The method comprises combining a test sample of AP with predetermined amount of a pseudo target (PT) and co-amplifying, to produce a collection of amplification products including an analyte amplicon if the sample contained AP and PT amplicon. The analyte amplicon is quantified without reference to the amount of PT amplicon and related to AP in the sample. The present sequence is a probe used in the rethod of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-182804/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nunomura K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GENP-) GEN-PROBE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JUL-1999; 99US-0145432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUL-2000: 2000WO-US20034
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Probe; HIV; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probe #2 used in a method for quantifying analyte polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-APR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF60545;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus type 1.
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 67 BP; 36 A; 9 C; 5 G; 17 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 39; 78pp; English
                                                                                                                                              0:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0:
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ID AAVS
                                 13-FEB-1998:
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SXCCCCCCCCCXPXPTTXX

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RESULT 15
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Best Local Similarity 100.0%; Prod. No. 35;
Matches 21: Conservative 0; Mismatches 0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lipid transfer protein coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Spinacia oleracea L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lipid transfer protein; membrane; liposome; drug carrier; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JUN-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA074410:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The lipid transfer protein encoded by this sequence may be used to alter the composition of lipid membranes. The lipid transfer protein may also be used to create new liposomes for use as divergences and new plants which have cells with altered membranes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI: 1994-322194/40.
P-PSDB: AAR63755.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-FEB-1993; 93JP-0035821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2: Page 2: 10pp: Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New lipid transfer protein and its gene \cdot useful for changing the composition of lipid membranes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NISB ) JAPAN TOBACCO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-FEB-1993:
                                                                                                                                                                                                                                                                             Serine protease: regulation: cell activity: viability: HELA2: ATC2: BCOM3; testisin: fertility; suppressor; testicular gorm cell cancer; seminoma; testic:specific expression; antitumour; sperm development infertility: mouse; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV59132 standard; DNA; 959
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 679 BP; 167 A; 134 C; 155 G; 223 T; 0 other;
                                      20-AUG-1998.
                                                                                    WD9836054-A1
                                                                                                                                                                                                                                    Hus sp.
                                                                                                                                                                                                                                                                                                                                                                                       Nucleotide sequence of murine HELA2
                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JAN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV59132;
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98WO-AU00085
                                                                                                                                                                       Location/Qualifiers
2..859
                                                                                                                           /*tag* a
/product= HELA2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ŖР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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PESULT 17
AAF22420C
XX
AAF224
XX
Human;
KW Human;
KW Human;
KW cerebr;
KW cerebr;
KW rheuma;
KW rheuma;
KW rheuma;
KW rocular
XW rheuma;
XW WO2000
XW WO2000
XW WO2000
XW WO2000
XW WO2000
XW WO2000
XW 19-OCT
XW WO2000
XW 19-OCT
XW WO2000
XW 19-OCT
XW 06-APR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match 3.6%; Score 21; DB 19; Best Local Similarity 100.0%; Pred. No. 34; Matches 21; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                             Human, secreted protein, diagnosis, immunosuppressive, antiarchritic:
antinheumatic, antiproliferative, cytostutic; andiant, vascinpic,
cerebroprotective; nootopic; neuroprotective; antibaceria; viricide;
fungicide; ophthalmological, gene theropy, antibaceria; viricide;
rheumatoid arthritis, hyperproliferative disorder; cardiac arrest;
cardiovascular disorder; cerebrovascular disorder; cerebral ischaenia;
anglogenesis; nervous system disorder; Arbalmer's disease; infection;
ocular disorder; corneal infection; wound healing; svin aging;
                                      06-APR-2000; 2000WO-US09071.
                                                                                                                                                                                                    WD200061629-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted protein gene 48 SEQ ID NO:58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF22420;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF22420 standard; cDNA; 1049 BP
                                                                                                                  19-OCT-2000.
                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 14: Fig 18A: 157pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-480768/41.
P-PSDB: AAW77301.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AMRA-) AMRAD OPERATIONS PTY LTD.
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13-FEB-1997; 97AU-0005101.
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AAF22373 to AAF22421 encode the human secreted proteins given in AAB53134
C to AAB53162. AAB53183 to AAB53231 represent more human secreted proteins
C and polypeptides homologous to them. Human secreted proteins have
C activities hased on the tissues and cells the genes are expressed in.
C Examples of activities include: immunosuppressive; antibacterial; vincide;
C inngicide; and pohthalmological. The polynucleotides and proteins can be
C used to prevent, treat or ameliorate a medical condition in e.g. humans,
mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They are
C also used in diagnosing a pathological condition or susceptibility to a
pathological condition. Disorders which are diagnosed or treated include
C pathological condition. Disorders which are diagnosed or treated include
C disorders e.g. neeplasms of the breast or liver, cardiouscular
C disorders e.g. neeplasms of the breast or liver, cardiouscular
C disorders e.g. contead by becteria, viruses and fungl and ocular
C disorders e.g. corneal infection. The polypeptides can also be used to
adjug due to sunburn, to maintain organs before transplantation, for
C supporting cell culture of primary tissues, to regenerate tissues and in
C supporting cell culture of primary tissues, to regenerate tissues and in
C chemotaxis. The polypeptides can also food additive or
C charcariza and Ana63133 represent sequences used in the exemplification of
C the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Covery Match 3.61; Score 21: DB 21: Length 1049: Best Local Similarity 100.0t: Pred. No. 3: Matches 21: Conservative 0: Mismatches 0: Indels
              Human: secreted protein: fusion protein: gene therapy; protein therapy; diagnosis: tissue: cancer; tumour: neurodegenerative disorder; leukaemia; dereloy-ental abnormality; foetal deficiency; blood; allerby; renal ds: imrune system; asthma; lymphocytic disease; brain; heptic; lymphoma; inflammaijon; ischaemic shock; Albeimer; disease; restenosis; Albe; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; tlymus; osteoprosis; arbhitis; lestis; lung; thyrodditis; thyroid disection; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
                                                                                                                                                                                                             Human secreted protein gene 75.
                                                                                                                                                                                                                                                         04-MAY-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                             AAX20485 standard; DNA; 1070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1049 BP; 305 A; 242 C; 212 G; 288 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1999; 99US-0128594.
20-JAN-2090: 2000US-0175931.
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(ROSE/) FOSEN C A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Caps
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AAC93427 standard:

ВP

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this sequence represents a nucleic acid molecule which encodes a secreted human protein. The gene number is given in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene can be used to generate fusion proteins by linking to the gene can be used to generate fusion proteins by linking to the gene can be fused protein as compared to the human protein only. The invention relates to 83 novel genes and their fragments (nucleic acid sequences) AXXXVIII:XXVIIII (2007) and their fragments (nucleic candiscide sequences) AXXVIII:XXVIIII (2007) and their fragments (nucleic conditions e.g. by protein or gene therapy. Also, pathological conditions e.g. by protein or gene therapy. Also, pathological conditions can be disposed by determining the amount of the new colymucleotides. Specific uses are described for each of the 95 colymucleotides. Specific uses are described for each of the 95 colymucleotides. Specific uses are described for each of the 95 colymucleotides based on which tissues they are most highly expressed in (see AXXVIII).
                                                                                                                                                          Ouery Match 3.5%; S. Best Local Similarity 100.0%; S. Matches 21; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Carter KC, E
19-MIC-1997
10-JTL-1997
10-JTL
                                                                                                                                                                                                                                                                                                                                                                Sequence 1070 BP; 298 A; 258 C; 198 G; 313 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 245; 312pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated human genes and the secreted polypeptides they encode useful for diagnosis and treatment of e.g. centers, neurological disorders, immune diseases, inflammation or blood disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI: 1999-153691/13.
P-PSDB: AAY00331.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-FEB-1999
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re PA, Rosen
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970S-0054211
970S-0054211
970S-0054211
970S-0054211
970S-0054214
970S-0054214
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970S-005595
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970S-005595
970S-005597
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CA,
                                                                                                                                                                     Score 21: DB
: Pred. No. 33:
0: Mismatches
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Puben SM, Shi Y, Wei
                                                                                                                                                                                                          DB 20;
. 33;
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                                                                                                                                                                                                                                     Length 1070:
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ei Y:
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Att State

Query Match Best Local Similarity

3.5%; Score 21; 100.0%; Pred. No

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21: Length 1164:

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CC human secreted proteins AAR53724 - AAR53777. Sequences AAR51778 -
CC human secreted proteins AAR51774 - AAR51777. Sequences AAR51778 -
CC AAR51825 represent alternative polypeptides encoded by the genes and
amino acid sequences to which they are homologous. The genes and proteins
cc have activities dependent on the itssues and cells in which they are
expressed. Examples of their activities include immunosuppressive;
cc antiarthritic; antirrhematic; antiroliferative; cytostatic cardiant;
visotropic; cerebroprotective; mostropic; neuroprotective; antibacterial;
cc virucide; funcicide; opthalmalogical; and vulnerary. The secreted
cc proteins, polymurisotides; antasynists and agonists may be useful in
cc treating, preventing and/or diagnosting diseases and disorders such as
autolamune diseases e.g. rhematoid arthritis, hyperproliferative
cc disorders e.g. neoplasms of the breast or liver, cardiovascular disorders
cc e.g. cardiac artest; cerebrovascular disorders e.g. cerebral ischaemia,
candiogenesis, nervous system disorders e.g. Alzheimer's diseases,
cc infections caused by bacteria, viruses and fund and ocular disorders
cc e.g. corneal infection. The polypeptides can also be used to aid wound
bealing and epithelial cell proliferation, to prevent skin aging due to
sunburn, to maintain organs before transplantation, for supporting cell
cc increase or decrease storage capabilities, fat content, lipid, protein,
cc polypeptides can also be used as a food additive or preservative to
components. Oligonucleotide AAC33413 - AAC3341 and peptide AAB51723 are
used in the isolation and characteris and other nutritional
cc polypucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human: secreted-protein: Immunosuppressive: antiarthritic; antirheumatic; antipressive; cyclostalic; cardiant; vasotropic; cerebroprotective; noctropic; neuroprotective; antibacterial; virucide; fungicide; opthalmalogical; vulnerary; autoimmune disease; rheumatoid arthritis; hyperproliferative disorders; cardiovascular disorder; cardiovascular disorder; cardiovascular disorder; cardiovascular disorder; altheumatoid stress; cerebrovascular disorder; nervous system disorder; Alzheimer's disease; ocular disorder; wound healing; skin aqing; ss.
  Sequence 1154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1: Page 413; 500pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid molecules encoding 48 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions used as food additives or preservatives
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI: 2000-619225/59.
P-PSDB: AAB51730.
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20-JAN-2000: 2000US-0177165
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(ROSE/) ROSEN C A.
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  BP:
284 A: 269 C: 310 G:
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301 T: 0 other
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ACC77607 to AAC78448 encode the human cancer associated proteins given the AAA3398 to AAB4429. The proteins can have activities based on the CLISBUES and cells the genes are expressed in Exemple of activities CL Include: cytostatic; proliferative; vulnerary; immunocipilator; cantidabetic; antiststmatic; antivienatic; antistrict; cantidabetic; antiststmatic; antivienatic; antistrict; cantidabetic; antiststmatic; antistrict; concurrence of the property; antibities; antistrict; cantidabetic; antistrict; antistrict; cantidabetic; antistrict; antistrict; cantidatetic; cantidatetic; antistrict; cantidatetic; cantidatetic; antistrict; cantidatetic; cantidat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ruman; cancer associated gene: cancer antigen; detection; cencer; diagnosis; cytostatic; proliferative; vulnerary; immunorodulator; antidibetic; antiashmatic; antirheumatic; antiarthritic; antiviral; antirhimatic; antipactic; antiviral; antirhimatic; antipactic; cardiant; dermatological; neuroprotective; thromblyric; cardiant; noctropic; antipacriatic; antianglogenic; gene therapy; inflammation; immune disorder; haematopojetic cell disorder; actormune disorder; antipacriatic; antianglogenic; gene therapy; inflammation; allergic reaction; graft versus host disease; organ rejection; haemostatic; thromblyric; cardiovascular disorder; infection; neurological disease; drug screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 1008: 2352pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated nucleic acids comprising sequences encoding reptides useful for treating or diagnosing e.g. cancer \cdot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-587533/55
P-PSDB; AAB43870.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human cancer associated gene sequence SEQ ID NO:473
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Sequence 1279 BP; 421 A; 225 C; 293

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337 T: 2

Covery with 1.59; Score 21: DB 22: Length 1279; Best Lett 15 indexity 100.01; Prod. No. 12: Patches 21: Conservative C: Mismatthes 0: Indels

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                                                                                                                                                                                            ARH3231 to AAH37195 and AAG73514 to AAG77788 represent human colon cancer associated nucleic acid molecules (N) and proteins (P), where the indecins are collectively known as colon cancer antiques. The colon cancer antiques that colon cancer antiques the colon cancer antiques the colon cancer antiques. The colon charity and vaccine production. Wand P may be used in the prevention, diagnosts and treatment of diseases associated with haspropriate P captrosts on. For example, N and P may be used to treat disorders cassociated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expression of P and distincially. N may be used to produce the colon cancer associated Ps. Committee the proteins or to supplement the patients own production of P. Additionally. N may be used to produce the colon cancer associated Ps. Committee the proteins. N and P can be used in the prevention, diagnosis and cancer colorectal carcinomas and cancers. Additional Colorectal carcinomas and cancers. Additional Colorectal carcinomas and cancers.
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AAH34592
ID AAH34582 standard: cDNA: 1279
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Rest Local Similarity 190.0%;
Matches 21: Conservative (
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                                                                present invention.
N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present for SEQ 'n NO:1027 to 1052. 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucloic acids encoding 4277 human colon cancer-associated polypeptides useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI: 2001-235357/24.
P-PSINA: AAG75177.
Sequince 1279 BP; 421 A; 225 C; 294 G; 337 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ruber SM. Barash SC. Birse CE.
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03-W-W-1999: 99US-0163280
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                                                                    Query Match 3.5%; Score 21; DB 17; Length 1372; Best Local Similarity 100.0%; Pred. No. 32; Matches 21; Conservative 0; Mismatches 0: Indels
The present sequence encodes an andothelial cell protein Cyactivated protein C receptor (EPPP). a type I transmembrane qlycoprofein. The protein binds with high affinity to both protein C (which plays a major role in blood coagulation) and activated protein C (Yd-30 nH) and is calcium dependent. It is a member of the CDL/MHC superfamily and has a role in regulating the inflammatory response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA encoding endothelial cell protein receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EPCR: endothelial cell protein C receptor: activated protein C:
blood coagulation: inflammatory response: inhibit; ss.
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                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 37-38; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                     Isolated endothelial cell protein C/activated protein C receptor used to inhibit inflammatory responses, screen for cpds. which alter receptor binding and, by blocking receptor binding, enhance inflammatory response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-AUG-1994; 94US-0289599
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                                                                                                                                                                          Sequence 1302 BP; 340 A; 325 C; 341 G; 296 T; 0 other;
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P-PSDB; AAR81453.
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CC vascular discase (especially cordiopulmonary bypass, unstable annima, respector in the conditions and the composed riversence of the receptor in the conditions.
                                                                                                                                       Courry Match 2.54: Score 21: DB 19: Length 1302: Best Local Sir:larity 100.07: Fred No. 32: Facthes 21: Conservative 0: Mismatches 0: Indels
Human; endo helial cell frotein receptor; inflammation; regulation; endothelial grotein C receptor; EPCR; coaquiation state; diabetes; major vascular condition; autoimmune disease; pre-eclampsia; cardiopulnomary bypass; unstable angina; restenosis; angioplasty; kidney disease; liver disease; as.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV53733:
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                                                                                                                                                                                                                                                                                                                                Sequence 13 2 BP: 340 A: 325 C: 341 G: 296 T: 0 other:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure: "olumns 21-22; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immuno-based detection of protein of receptor - useful in the diagnosis of inflammatory and coaquilation states and disorders associated with damage to endothelium and large blood vessel disease
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P-PSDB: AAW 4546.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Esmon CT, Furosawa S. Stearns-kurosawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (OKLA-) OKITHOMA MEDICAL RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27 - JUN-1997 -
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25..741
/*tag= a
/product= "ECFR protein"
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                                                                                                                                                           Gaps
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RESULT 24
AAA07076
ID AAA070
Chamban endothelial protein C receptor (EPG). The invention relates to a modified form of EPGR, in which the C-terminal Cys of the membrane-to a condition form is not paintoylated, or is otherwise replaced by a different camino acid; to a soluble splice wariant of EPGR, which lacks a transmembrane domain; and to a EPGP in which the transmembrane domain of the membrane-bound form has been removed by proteolytic cleavage. Splicing of exon III to exon IV (which encodes the transmembrane domain) the EPGR RNA transcript results in an mRNA encoding the soluble form cludes intron III sequences. The intron III sequence includes an inframe termination codon: translation of intron III containing mRNA cresults in an EPCR protein with a variant sequence (ANSH177) immediately CC C-terminal of glycine 201. EPCR binds both protein C and its activated code form, activated protein C (APC), with similar affinity in the presence of calcium. The membrane-bound form facilitates protein C activation by competing with membrane-bound EPCR for protein C complex on cell surfaces. In contrast, the soluble form inhibits protein C calcium. The membrane-bound Grow facilitates protein C activation of protein C to a thrombin/thrombomodulin activation complex on calcium. The competing with membrane-bound EPCR for protein C. The calcium in of blood coagulation. Pattents deficient in protein C calcium. The competing with membrane-bound EPCR for protein C. The calcium in of blood coagulation. Pattents deficient in protein C. In the recein C. usually exhibit life-threatening thrombotic complications in infancy.
               Claim 4; Columns 21-22; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modified endothelial protein C receptor, useful as diagnostic marker of e.g. Inflammation or autoimmune diseases, is a soluble or alternatively spliced form of the receptor.
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P-PSDB; AAT81476.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Endothelial protein C receptor: EPCR; membrane bound: splite variant: activated protein C: APC: coequiation: inflammation: regulation: endothelial cell: autolmmune disease: vascular damage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JUL-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA07076 standard; cDNA; 1302
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25.74
/*tsg- a
/*roduct- *Human membrane-bound EPCR splice variant*
25.69
25.69
/*tsg- b
/*tsg- b
/*tsg- b
/*tsg- b
/*tsg- c
/*product- *Mature human membrane-bound EPCP*
627.628
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/note- "Exon III-exon IV junction"
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ARESULT 25
AAZ98076
AX AAZ9807
AX AAZ980
AX Human:
AW Human:
AW Human:
AW Human:
AW Hotelal
AW Hotelal
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15-JUL-1998;
15-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human: secreted protein: diagnosis: cytostatic: immunosuppressive: antiinijammatory; nootropic; neuroprotective; antialiergic; cancer: tumour; neurodegenerative disorder: developmental ahordmality; allery; foetal deficiency; blood disorder: jamune system disorder; arthritis; autoimmune disease; hepatic disease; renal disease; inflammation; Alzhe'mer's disease; behavioural disorder; schizophrenia; osteoporosis; infection; hills; spinal cord injury; transplant rejection; diabetes; asthma: sepsis; acne: psorlasis; catdiovascular disorder; reproductive disorder; gastronivestinal disorder; respiratory disorder; metabolic disorder; food additive; preservative; ss.
                                                                                                       Claim 1; Page 346-347; 494pp; English.
                                                                                                                                                                 New isolated human genes, useful \tau diagnosis and treatment cancers, neurological or blood distrders -
                                                                                                                                                                                                                                                                         WFI: 2000-151128/14.
P-PSDB: AAY87123.
                                                                                                                                                                                                                                                                                                                                                                           Ruben SM. Komatsoulis G. Duan PP. Poson CA. Moore FA.
Laflour DW. Ebner R. Olsen HS. Frewer LA. Florence KA.
Mucenski M. Endress GA. Soppet PP:
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The polynucleotide sequences given in \Delta AZ98017 to \Delta AZ98108 encode the human secreted proteins given in \Delta \Delta Y87064 to \Delta AY87223. Human secreted
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98US-0092922.
98US-0092956.
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PESULT 26
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communosuppressive antinflammatory; nootropic: neuropycontice; and antiallergic. The polymerides and their corresponding secretal polymerides are useful for preventing, treating or arelibration effect can be diagnosed by determining the amount of the new polymerides in a graphe or by determining the presence of mutations in the new polymerides in a polymerides determining the protein s and their polymerides in a polymeride coulder. Human secreted protein s and their polymerides in a polymeride coulder. Human secreted protein s and their polymerides can be used for developing products for the diagnosis or training the gamer, tumburs, neurodegenerative disorders, developmental abnormalities and feel clencies, blood disorders, developmental abnormalities and feel clencies, blood disorders, developmental abnormalities and altergies. Alzhener's disease, behavioural disorders, eshiration, allergies, alzhener's disorders, and also be used as food additives of the development and allergies allergies
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Best Local Similarity 100.0%;
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human: secreted protein: proliferative disorder: cancer: "nour: asthra; foetal abnormality; developmental abnormality; haveacopolic disorier; for immune system disorder: AIDS; autoimmune disease; thermand at mirrier parkinson's disease; cognitive disorder; schizophrenia: skin disorder: psorlasis: sepsis; diabetes: atheroscierosis; cardiovascular disorder: inflammation; neurological disorder: Alzheimer's disease; icol stillier; angiogenic disorder; tidney disorder; alzheimer's disease; icol stillier; pregnancy-related disorder: endocrine disorder: Infection: wound bealting gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             sig_peptide
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                                                                                         19-JUL-2001.
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                                                                                                                                                                                                                                                                                                                                                              /*tag= c
/product= "Mature human secreted protein"
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Pred. No. 32;
0: Mismatches 0: Indels
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12-JAN-2001: 2001WO-US90311

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CC The secreted proteins and their genes are useful for proteins they encode. The secreted proteins and their genes are useful for proteins or gene therapy to a maniforation redical conditions, e.g., by protein or gene therapy to a maniforation redical conditions can be diagnosed by determining the amount of the CC the new genes, specific uses are described for each of the 71 games. In the seed on the tissues in which they are most highly expressed and include CC developing products for the diagnosis or treatment a brother title. CC the new genes, specific uses are described for each of the 71 games. In CC developing products for the diagnosis or treatment a brother it is considered. CC developing products for the diagnosis or treatment a brother titles. CC developing products of the 1mmune system, AIDS, autolomous discorders, cancer turours, forein and developmental abnormal titles. CC discases (e.g., rebumated) at this is an indevelopment of the 1mmune system, AIDS, autolomous discorders, cancer turours, forein and development of the 1mmune system, AIDS, autolomous discorders, cancer turours, forein and architecture, although stin discorders (e.g., AIDS, autolomous discorders, cancer (e.g., AIDS, autolomous discorders), reportants, septis, althoryers, gastrointestinal discorders. CC analogonit discorders kidney discorders, sastrointestinal discorders, and indevelopment of the analogonit discorders and considers, cardiovascular discorders. The proteins can also be used to all wound healthman of printering tissues, to regular discorders and include the proteins are also be used to all wound healthman of printering tissues. The proteins can also be used to all wound healthman, to maintain organic printers, and in chemotaxits, and can be used as a food additive or printering of the invention of the invention can be used as a food additive or printers, and in chemotaxits, 
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Best Local Similarity 100.0%; F
Matches 21; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytokine: zsig 81; wound healing: proliferation; differentiation migration: metabolism; ss.
                                                                                                                                                                                 Mouse ortholog of zsig81 ENA.
                                                                                                                                                                                                                                                        19-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                         AAC91065;
                                                                                                                                                                                                                                                                                                                                                                                                           AACG1045 standard: DNA: 1547
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F-FSDB: AAE05100.
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Lafleur DW, Olsen HS, Brewer LA, Florence KA,
Endress GA, Muscenski M, Fhner B;
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Young PE,
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, Soppet DR;
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RESULT 28
AAD08465
AD08465 standard; cDNA: 1549
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Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                      Human; secreted protein, prolife in the disorder; cencer; turour; asthra; foetal abnormality; haematopoietic disorder; immune system disorder; AIDS; autoimmune disease; rheumatoid arthritis; psorlasis; sepsis; dabetes; atherosclerosis, cardiovascular disorder; psorlasis; sepsis; dabetes; atherosclerosis, cardiovascular disorder; andiogenic disorder; kiney disorder; Albeimer's disease; food additive; andiogenic disorder; kiney disorder; alteigner's disease; food additive; pregnancy related disorder; endocrine disorder; infection, cound healing; cell culture; chemotaxis; vulnerary; binding partner identification; gene therapy; chromosome 20; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-MAT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted protein-encoding gene 17 cDNA clone HHENZ15, SEQ ID NO:72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-AUG-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to zsig@l and fragments thereof. The invention is useful for promoting would healing, for modulating the proliferation, differentiation, migration or metabolism of responsive cell types that includes both primary and cultured cell lines, and for stimulating the proliferation of cells expressing markers associated with dendritic lineage cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-2000; 2000WO-US15002
                                                                   WO200134643-A1.
                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1547 BP; 428 A; 375 C; 357 G; 387 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure: Page 95-96; 109pp: English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New rsigal polypeptides and polypucleotides useful for e.g. gromoting would healing, or in diagnosing or treating disorders associated with cell loss or abnormal cell proliferation, such as cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Piddington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200073459-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ch 3.5%; Score 21; DB 22; Length 1547; Similarity 100.0%; Pred. No. 31; 121; Conservative 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     West JR, Holly RD,
                                                                                                                     Location/Qualifiers
795.809
/*tag a /*product* *Human secreted protein precursor*
/note* *Does not include start codon*
/partial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Burkhead
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0:
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fructose-6-phosphate 1-phosphotransferase enzyme;

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                                                                                                                                                                                                                                                                                                                                Query Match 3.5%;
Best Local Similarity 100.0%;
Matches 21; Conservative
                                                                                                                                                                                                                                          Oryza sativa pPEK-OSI dene
                                         12-001-1995 (first entry)
                                                                                                                                  AACRES84 Standard:
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1549 BP; 411 A; 302 C; 305 G: 529 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acids encoding 24 human secreted polypeptides, useful for preventing, diagnosing and/or treating e.g. Gaucher's disease, Alzheimer's disease, Scimiter syndrome, Crentzfeldt-Jacob disease, Scimiter syndrome, Crentzfeldt-Jacob disease, diabetes mellitus and multiple sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI: 2001-374441/39
P-PSDB: AAE04160.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ruben SM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Komateoulis GA.
                                                                                                                                  CDNA to mRNA: 1624
                                                                                                                                                                                                                                                                                                                                   Score 21: DB:
Pred. No. 31:
0: Mismatches
                                                                                                                                                                                                                                                                                                                                0:
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                                                                                                                                                                                                                                                                                                                                                                             22;
                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                             Length 1549:
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RESULT 30
AAI58747
ID AAI5874
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1600 atatttaaaaaaaaaaaaa 1620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequences (AAQ85982-6) represent the genes encoding a novel ATP-dependent fructose-6-phosphate 1-phosphotransferase enzyme (EC 2.7.1.11; PEK) from a range of plants. This is the sequence of the OTYZA SATIVA (Fice) gene, pPFK-OSI, as given in the specification. Plants transformed with these genes can express the enzyme. The transformed plants can produce varieties that have altered sugar moniton on storage at low temperatures.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 8; Page 46-49; 79pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA coding for fructose-6-phosphate 1-phosphotransferase - of plant origin, for prodm. of transformant plant cells with altered sugar content
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hiyoshi T. Kasaoka K, Mine T. Page MJA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-AUG-1993; 93JP-0226454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-AUG-1994;
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Raphanus sativus: rlaveria brownii: primer: expression vector:
Agrobacterium tumefaciens: sugar; storage: temperature: ds.
                                                                                      Human, moduropic, immunosuppressant; cytostatic, gene therapy; car er; peripheral netwous system, neuropathy; central netwous system; CNS-Alibelmer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral scierosis; Shy Drager Syndrone; chemo-dattic; chemo-kinetic; thrombolytic; drug screening; arthritis; inflammatic;
                                                                                                                                                                                                                                                                                                                         22-0CT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                    AA158747;
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAI58747 standard: cDNA: 1753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1624 BP; 480 A; 302 C; 406 G; 436 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NISB ) JAPAN TOBACCO INC.
                                                                                                                                                                                                                                                              Human polynucleotide SEO ID NO 950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       570 atatttaaaaaaaaaaaaa 590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-098757/13.
P-PSDB; AAR71581.
Homo sapiens.
                                                              leukaemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94WO-JP01352
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/transl_except* seq: AAT, a.a.:Ile
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/product- fructose-6-phosphate 1-phosphotransferase
390..392
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3..1412
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RESULI 31
AAH48294
ID AAH482
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.6%; So Best Local Similarity 100.0%; F Matches 21; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21 - JAN - 2000;
25 - APR - 2000;
09 - JUL - 2000;
19 - JUL - 2000;
03 - AUG - 2000;
19 - OCT - 2000;
29 - NOV - 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1: SEQ ID NO 950: 10078pp: English.
Honeybee; alpha-qlucosidase; ds
                                                   Honeybre alpha-glucosidase coding sequence.
                                                                                                      25-SEP-2001 (first entry)
                                                                                                                                                      AAH4 B 294 :
                                                                                                                                                                                                          AAH482n4 standard: DNA: 1994 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1763 BP; 527 A; 327 C; 399 G; 510 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel nucleic acids and polypeptides, useful for treating disorders such as central nervous system injuries -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI: 2001-442253/47.
P-PSDB: AAM39591.
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Wang J, Wang Z, Wehrman T, Xu C, Xue AJ,
Zhao QA, Zhou P, Goodrich R, Drmanac RT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HYSE.) HYSEQ INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-JUL-2001.
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2000US-0552317.

2000US-05598042.

2000US-0620312.

2000US-0620312.

2000US-0652191.

2000US-0693035.

2000US-0693035.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 21; DB 22; Length 1763; Fred. No. 31: 0: Mismatches 0: Indels
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Zhang J;
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RESULT 32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim la; 116–117; 188pp; German
                                   Human nucleic acid sequences and protein products from tissue, useful for breast cancer therapy -
                                                                                                                                                                                                                                   20-MAR-1998;
                                                                                                                                                                                                                                                                       20-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                 DE19813839-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                 Expressed sequence tag: ESI; human; breast; cancer: gene therapy;
treatment; tumour: cytostatic: medicament; ss.
                                                                                                                                                       Specht T, Hinzmann B, Schmitt A, Pilarsky C.
                                                                                                                                                                                                                                                                                                           23-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                     Romo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human breast tumour-associated EST 37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-DEC-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ33647 standard; cDNA; 2281 EP
                                                                                                                                                                                          (META-) METAGEN GES GENOMFORSCHUNG MBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1994 BP; 753 A; 310 C; 357 G; 574 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides the protein and coding sequences of the nomeybee alpha-glucosidase. The gene can be used for the preparation of Apis mellifer a alpha-glucosidase I. The present sequence is the coding sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 6-9; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Apis mellifera alpha-glucosidase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-460212/50.
P-PSDB; AAG64875.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-SEP-1999; 99JP-0245862.
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                                                                                                                                                                                                                                 98DE-1013839.
                                                                                                                                                                                                                                                                       98DE-1013839.
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/product= *alpha-glucosidase*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualiflers
13..1779
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22;
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                                                          tumor breast
                                                                                                                                                       Rosentahl A:
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Ouery Match
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RESULT 33
AA090652
ID AA09065
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Best Local S
Matches 21
Sequence 3133 BP: 718 A: 918 C: 922 G: 575 T: 0 other:
                                        Novel FPH: related PTK Ceké CDNA clones (AA09052) were isolated from chick erbiyo and embryonic balan CDNA, libraries in phage lambda (LDNA). CDNA clones (ABC) related to rat EDNA). Ceké CROSSON (CEKÉ (ACC) and CENA) (AAR75704). Ceké Clanscripts were found in Ceké CAAP7572) and CENA (AAR75709). Ceké Clanscripts were found in 10-day embryos and in adult brain, lung, heart and skeletal muscle.
                                                                                                                                                                                      Disclosure: Page 37-41: 129pp: English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI: 1995-215256/28.
P-PSDB: AAR75704.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09515375-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gallus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cek6: Eph: protein tyrosine-kinase; PTK; cancer; diagnosis; prognosis: ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eph-related tyrosine kinase CEK6 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ90652;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention devitbes movel human nucleic acid sequences from tumor breast tissue which have cytostatic activity. The nucleic acid sequences can be used to produce and isolate full-length gene sequences. They can be used to express proteins, which can be used as tools to find an activity against heast cancer. The sequences can be used in sense or antisense form. They are especially useful for medicaments for gene therapy to truck hreast cancer. Ac233(1):246(7) represents expressed sequence tags described in the method of the invention.
                                                                                                                                                                                                                                                 Eph-related protein tyrosine kinase(s) - for monitoring and diagnosing
                                                                                                                                                                                                                                                                                                                                                                     Pasquale EB, sajjadi FG:
                                                                                                                                                                                                                                                                                                                                                                                                                        (LJOL-) LA JOLLA CANCER RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-NOV-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA090552 standard: cDNA: 3133 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2281 BP; 501 A: 498 C; 494 G; 688 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           atch 3.6%; Score 21; DB 20; Length 2281; Cal Similarity 100.0%; Prod. No. 30; Cal Similarity Conservative 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9305-0162809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94WO-US10140
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421..2859
/*tag- b
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3..419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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3.6%:

Score 21;

DB 15;

Length 3133;

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RESULT 35
AAX39661/c
                                                                                               Query Match
Best Local S
Matches 21
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Matches 21; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is the coding sequence for portine CD29 protein. CD29 contains rpitopes Gal-alpha-(1.3)-Gal, which are recognised by genoreactive human antibodies, leading to hyperacute rejection of xenograffs. The present invention relates to CD29 proteins, which have an identical, reduced or zero expression of epitope Gal-alpha-(1.3)-Gal, which can be used to reduce or prevent hyperacute rejection associated with xenotronsplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Porcine; CD29; immunosuppressive; immunomodulatory; epitope Gal-alpha-(1.3)-Gal; xenotransplantation; xenograft; ss.
                                                                                                                                                                                                                                                                                     Claim 4; Page 38-39; 46pp; Spanish.
                                                                                                                                                                                                                                                                                                              Porcine CD29 protein and related DNA, antibodies to prevent graft rejection useful as graft donors -
                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-273559/28.
P-PSDB; AAB84751.
                                                                                                                                                                                                                                                                                                                                                                                                    Garrido Pavon JJ, Llanes Ruiz D, Jimenez Marin AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                       (BIOV-) BIOVET-UCO SL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-OCT-1999; 99ES-0002193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-OCT-2000; 2000WO-ES00374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Porcine CD29 protein coding sequence.
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                                                                                                                                                             Sequence 3831 BP; 1114 A; 703 C; 920 G; 1094 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   34
                                                                                             tch 3.6%; Score 21; DB 22; Length 3831: al Similarity 100.0%; Pred. No. 28; 21; Conservative 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
241.7637
/ttage a
/product * Porcine CD49*
/transl_except * (pos:1027..1035,aa:Ser-Leu-Ile)*
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); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                 Babancho Medina
                                                                                                                                                                                                                                                                                                                          useful for removing xenoreactive and to prepare transgenic animals
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                                                                                                 Gaps
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                                    AAMSOS30/C
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                                                                                                                                     Overy Match 3.5%; Sometime 100.0%; From Matches 21; Conservative 0;
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17-JUL-1997;
10-OCT-1997;
10-OCT-1997;
10-OCT-1997;
11-OCT-1997;
                                                                                     Claim 67: Page 500-502: 787pp; English.
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   AAN50530;
                       AAN50530 standard: DNA; 5750 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI: 1999-132448/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (LUDW-) LUDWIG INST CANCER RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Renal cancer associated gene.
                                                                                                                                                                                                                            lung cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0102322.
97US-0896164.
97US-0061599.
97US-0061765.
97US-0948705.
97GB-0021697.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98WO-US14679
                                                                                                                                         Score 21: DB:
; Pred. No. 27:
0; Mismatches
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The invention relates to a method for diagnosing a disorder characterised by expression of a human cancer associated antigen precursor coded for by a nucleic acid molecule (AMA). The method comprises: (a) contacting a biological sample isolated from a subject with an agent that specifically binds to the NAM, an expression product or a fragment of an expression product complexed with an HLA molecule; and (b) determining the interaction between the agent and the NAM or the expression product as determination of the disorder. The products and methods can be used in the diagnosis, monitoring, research, or treatment of conditions characterised by the expression of various cancer associated antigens. The invention provides nucleic acid sequences and encoded polypeptides which are cancer associated antigen precursors expressed in human breast cancer, remail cancer, colon cancer, gastric cancer, prostate cancer and und cancer.
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breast cancer: colon cencer: gastric cancer: renal cancer: lung cancer:
prostate cancer: ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen Y, Gout I, Gure A, O'Hare M, Obata Y, Old LJ:
Pfreundschuh M. Sahin U, Scanlan MJ, Stockert E:
Tureci O:
                                                                                                                                                                                                                                Sequence 5305 BP; 1857 A: 973 C: 1076 G; 1390 T: 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          have isolated cancer associated nucleic acids and polypeptides -
isolated using sera from cancer patients, used to develop products
for the diagnosis, monitoring or treatment of cancers
                                                                             20; Length 5306;
0:
Gaps
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AAC71811
AAC778
XX
AAC778
AC AAC778

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22-FEB-1984;
21-FEB-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence encoding the P195 protein of Plasmodium falciparum (AAM5030) and a peptide comprising at least one of its epitopes (see AAP5077) are claimed. Also claimed is a vaccine for inducing immunity to malaria comprising the novel peptide or P195 or a peptide comprising at least one epitope when derived from the new DNA sequence, together with a carrier.
Human: cancer associated gene: cancer antigen: detection: cancer: diagnosis: cytostatic; proliferative: winerary: immunocollator: antidiabetic; antiasthmatic; antirhematic; antiarthricic; antiviral: antiinflammatory; antityloid; antiallergic; antibacterial; cardiant; dermatological; neuroprotective; thromblytic; coagulant; novtopic; wasotropic; antipsociatic; antiandiogenic; gene therapy; inflammation: immunocdisorder; hemenocopietic cell disorder; autoimmunocollatic; antiandiogenic; gene therapy; inflammation: allergic reaction; graft versus host disease; organ rejection:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC77811;
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                                                                                                                                                                                                                                                                                                                                                                                                                          Human cancer associated gene sequence SEO ID NO:205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC77811 standard;
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P-PSDB; AAP50777.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-SEP-1991 (first entry)
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84GB-0004692.
85GB-0004429.
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216..5179
/*tag= a
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RESULT 38
ANX/51565
ANX/51565
ID AAX/51565
XX ACC AAX/51565:
XX O5-AUG-1999 (first entry)
DI 05-AUG-1999 (first entry)
XX Pneumocystis carinii lysyl-ti
XX Anthony Ant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC ANAC77607 to ANG/8448 encode the human cancer associated proteins given CI in ANAH 3138 to ANAH2789. The proteins can have activities based on the Cissues and colls the genes are expressed in Example of activities CI include: cytostatic; proliferative; vulnerary; immunemedulator; cantidabetur; antiasthantic; antiferative; multaribities: CC autidabetur; antiasthantic; antiferative; antiaribities: coagulant; coagu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Owery Match 21: Lenoth 5830: Score 21: DB 21: Lenoth 5830: Best Local Similarity 100.0%; Pred No. 26: Matches 21: Conservative 0: Mismatches 0: Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-MAR-2000; 2000WO-US05882.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-587533/55.
P-PSDB; AAB43602.
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                                                     pneumorystis carinii lysyl-tana synthetase: tyrosyl-tana synthetase:
aminoacyl-tFNA synthetase: pneumonia: AIDS: immuno-compromised:
Acquired Immune Driftclency Syndrome: detection: PCR primer: ss.
                                                                                                                                                                                                                           Pneumocystis carinii lysyl-tRNA synthetase PCR primer #29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5830 BP; 1558 A: 1417 C; 1409 G; 1439 T; 7 other;
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us-09-248-178-67.rng

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RESULT 39
AAX76360/C
ID AAX76360 standard; DNA; 40 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes Pneumocystis carinii (P. carinii)

Caninoacyl-RNA synthetase enzymes. The nucleic acids encoding aninoacyl-
CRNA synthetase enzymes. The nucleic acids encoding aninoacyl-
CRNA transpares may be used to produce expression vectors and

Chost cells for the recombinant production of pneumocystis aminoacyl-
CRNA

Synthetase. The proteins may then be used in other procedures such as

CRNA

C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 3.4%; So
Best Local Similarity 100.0%; I
Matches 20; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acids encoding Pneumocystis carinii aminoacyi-tRNA synthetase enzymes useful for detecting similar sequences in samples and in the study and treatment of pneumonia in Acquired I-mune Deficiency Syndrome patients
15-JUN-1999.
                                               US5912140-A.
                                                                                        Synthetic.
Pneumocystis carinii.
                                                                                                                                                             Pme:mocystis cartinli, lysyl-tENA synthetase: tyrosyl-tENA synthetase: manionacyl-tENA synthetase: pneumonia: AIDS; immuno-cc-promited: Acquired Immune Deficiency Syndrome: detection: PCR primer: ss.
                                                                                                                                                                                                                                                           Pneumocystis carinii lysyl-tRNA synthetase PCR primer 31.
                                                                                                                                                                                                                                                                                                          05-AUG-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                         AAX76360;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 17; Column 47; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Politis-Virk KI, Quinn CL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-APR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 35 BP; 7 A; 5 C; 2 G; 21 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-357196/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CUBI-) CUBIST PHARM INC.
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; Pred. No. 1.2
0; Mismatches
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. 1.2e+02;
ches 0;
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Q 8

95WO-GB01987

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RESULT 40
AAT17027/c
ID AAT17027 standard: DNA: 42
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annoacyi-ERNA synthetase enzymes. The nucleic acids encoding aminoacyi-
ERNA synthetase enzymes may be used to produce expression vectors and
thost cells for the recombinant production of Pneumocystis aminoacyi-ERNA
Synthetases. The proteins may then be used in other procedures such as
c separating amino acids from samples or as antigens in the production of
antibodies. The nucleic acids may also be used to produce tester cell
c strains (which contain the nucleic acids) which may be used to test
c candidate drugs (e.g. tRNA synthetase inhibitors) for the treatment of
c disorder associated with P. carinii such as pneumonia which is a common
c complication for acquired imanue beficiency Syndrome (AIDS) patients and
c other immuno-compromised individuals. Additionally, they may also be
used to detect and isolate related DNAs in sample (i.e. they can be used
c as probes). The present segmence represents a PCR primer for a
c present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 3.4%: Sometime State Sest Local Similarity 100.0%; I Matches 20; Conservative 0;
29-FEB-1995
                                                         K29606187-A1
                                                                                                                                                                                 misc_feature
                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                  Human; mitochondrial DNA, heavy chain; primer; polymorphism;
identification; discrete single nucleotige; bases; method; screen;
genetic disease; DNA typing; forensic testing; microorganisms; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mitochondrial DNA heavy chain primer H16070.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94-OCT-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40 BP; 8 A; 6 C; 4 G; 22 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 16; Column 44; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acids encoding Pneumocystis carinii aminoacyl-tRNA, synthetase enzymes useful for detecting similar sequences in samples and in the study and treatment of pneumonia in Acquired immune Deficiency Syndrome patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI: 1999-357196/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-APR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-0415593
                                                                                               /*tag* a
/note= "5'-polyT tail. opt. polyA"
                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 20; DB 20; 1
Pred. No. 1.2e+02;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT 41
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A sample is mixed with primers (P) that are complementary to the characterised base sequence so that they annual to the nb, at classically types of chain terminator each having a characteristic fluorescent label, and a chain extending enzyme so that the extended P are then sept. according to size or charge, and the extended P are then sept. according to size or charge, and the screen for genetic diseases associated with single nb, and for DNA typing, e.g. in forensic testing, it may also be used to screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb, and for DNA concerns the service of the screen for genetic diseases associated with single nb.
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Best Local Similarity 100.0%;
Matches 20; Conservative (
(PHYL-) PHYLOS INC
                                                                                        02-DEC-1998;
                                                                                                                                                                02-DEC-1999;
                                                                                                                                                                                                                                                                                                                                      WO200032823-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA-protein fusion; RNA-protein fusion; stable; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Model mPNA #3 used to demonstrate DNA-protein fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JAN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA61354 standard; mPNA; 51 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 42 BP; 7 A; 2 C; 8 G; 25 T; 0 other:
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by annealing primer adjacent to the base and extension with
fluorescently labelled chain terminator, for simultaneous screening
of genetic diseases, or for forensic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI: 1995-151390/15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SOLI-) SOLICITOR AFFAIRS OF HM TPEASURY.
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                                                                                        9805-0110549
                                                                                                                                                                99WO-US28472
                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
6..28
/*tag* a
/bound_moiety* *Linker 7*
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AAV25706
ID AAV25706 standard: cDNA; 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to methods of covalently tagging proteins with their encoding DNA sequences. Several methods may be used to create these DNA-protein fusions. In owneral, the methods involve linking a corpiner to an RNA molecule, then translating the RNA to produce a protein which is bound to the primer. The RNA is then reverse transcribed to complete the translating the RNA to produce a protein corpine as used to demonstrate DNA-protein fusion. The present sequence is model mRNA is contained to that was used to demonstrate DNA-protein fusion formation. This cosequence was used as a substrate for the binding of linker 7 (see CC AAA61351). The DNA-protein fusions are useful in binding and molecular corporation assays that involve biological materials containing continuous seases. The covalent bond between the protein and the encoding CDNA makes the fusion more stable than RNA-protein fusions. This means conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 3.4%;
Best Local Similarity 80.0%;
Matches 16; Conservative
           Compositions containing isolated proteins encoded by specific nucleic acids - useful e.g. as anti-inflammatory, immune stimulant or suppressor, etc.
                                                                                                                    Jacobs K, Lawallie ER, McCoy JM, Merberg D, Spaulding V, Treacy M:
                                                                                                                                                                                                            04 - OCT - 1996;
                                                                                                                                                                                                                                                03-CCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                       Secreted protein, human, marker, cytokine; immune stimulant; suppressor; autofommune disease; regulator; activin; inhibin; inhibitor; chemotactic; chemokinetic; harmostatic; throrkocyte; tumbur; antilinfiammatory; ds.
                                                                                                                                                                        (GEMY ) GENETICS INST INC.
                                                                                                                                                                                                                                                                                 09-APR-1998.
                                                                                                                                                                                                                                                                                                                     W09814470-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-SEP-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 51 BP; 26 A; 7 C; 5 G; 7 U; 5 other:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Separating a DNA-protein (usion useful for selecting a desired protein
or the encoding DNA, comprises covalently linking proteins with their
encoding DNA sequences
                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human novel secreted protein clone BF290_li 3'~end DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure: Fig 14: 51pp; Enalish.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lohse P. Kurz M.
                                                                                                                                                                                                            9605-0725885
                                                                                                                                                                                                                                                97WD-US18032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wagner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           : Score 20: DB 21;
: Pred. No. 1.2e+02;
4: Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lenath 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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Query Match 3,4%; So
Best Local Similarity 100.0%; F
Matches 20; Conservative 0;
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The present invention relates to methods of covalently tasking proteins with their encoding DNA sequences. Several methods may be used to create these DNA protein fusions. In general, the methods involve linking a primer to an RNA molecule, then translating the RNA too produce a protein which is bound to the primer. The PNA is then reverse transcribed to give a DNA-protein fusion. The present sequence is model PNA substrate in that was used to demonstrate DNA protein fusion formation. The DNA-protein fusions are useful in binding and molecular recognition assays that involve biological materials containing ribonucleases. The covalent bond between the protein and the encoding PNA makes the fusion more stable than PNA-protein fusions. This means selection experiments need not be limited to extremely mild reaction conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-DEC-1998; 9805-0110549
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA-protein fusion; RNA-protein fusion; stable; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 15; 5lpp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Generating a DNA-protein fusion useful for selecting a desired protein or the encoding DNA, comprises covalently linking proteins with their encoding DNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lohse P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PHYL-) PHYLOS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Model RNA substrate #3 used to demonstrate DNA-protein fusion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 79:
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Best Local S
Matches 16
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                                                                                                                                                                Sequences AAA31040-A32093 represent nov-1 plant microsatellite sequences and associated flanking species. The sequences comprise a central core repeat sequence, especially selected firm the sequences AAA32094 A32096 with left and right flanking sequences. The polynuciections sequences can be used in the detection of DNA polynophisms, in genome mapping, in positional cloning of genes, in variety identification and in evaluation of genetic variability within and between plant tissues, populations, cultivars, species and species groups. They may also be used to design hybridization probes for colligonucleotide ingerprinting and livary screening and to design primers for microsatellite primer for microsatellite primer for microsatellite primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 115 BP; 48 A; 23 C; 23 G; 21 U; 0 other;
                                                                                                                                       Sequence 242 BF; 106 A; 23 C; 70 G; 43 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 337; 392pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                New plant microsatellite markers and associated flanking species for
the detection of polymorphic genetic markers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Havukkala IJ. Bloksberg LN. Glenn M:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENE-) GENESIS RES & DEV CORP LID & FLITCHER (FLET-) FLETCHER CHALLENGE FORESTS LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plant microsatellite sequence; core regeat sequence; detection; probes
DNA polymorphism; genore.mapping; physical mapping; finecrprintings;
variety identification; cenetic variability evaluation; primer; ss.
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91 uauuuaaaaaaaaaaaaa 110
tch 3.4%; al Similarity 100.0%; 20: Conservative
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15; Conservative
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RESULT 45
AAZ08301
ID AAZ08301 standard; cDNA; 278 BP.
Query Match 3.4%; Score 20; DB 20; Length 278; Best Local Similarity 100.0%; Pred. No. 93; Natches 20; Conservative 0; Hismatches 0; Indels
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28-JAN-1998;
28-JAN-1998;
18-MAR-1998;
18-MAR-1998;
23-JUL-1998;
23-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes lung tumour specific polynucleotides and tumour antigens. AAC07144 to AAC07245 and AAC08257 represent specifically claimed polynucleotides, and AAV23485 to AAV23571 represent amino acid sequences from the present invention. The lung tumour specific polynucleotides and polypeptides can be used in pharmaceutical compositions and vaccines to inhibit the development of lung cancer. They can also be used to detect lung cancer in a patient. Probes and antibodies derived from the lung tumour sequences are useful in detection of lung cancer.
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                                                                                                                                                                                   Human lung tumour protein SAL-109 5' cDNA sequence.
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Maximum DB seq length: 2000000000
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Gapop 60.0 , Gapext 60.0
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Copyright (c) 1993 - 2000 Compugen Ltd
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## ALIGNMENTS

DNA encoding human breast tumour protein immunogenic fragment.

Breast tumour protein; immunogenic fragment; vaccine; detection; breast cancer development; therapy; ss.

New preast tumour protein genes used, in vaccines for immunotherapy, or for diagnosis of breast cancer

3; Page 60; 70pp; English

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Best Local Similarity 100.0%;
Matches 1079; Conservative 0
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RESULT 2
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ID AAC79438 Standard; c
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AC AAC79438;
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CDNA sequence of human breast tumour clone 1015D11

07-FEB-2001 (first entry)

Human; breast tumour antigen; cytostatic; immunotherapy; breast cancer; vaccine; ss.

WD200061756-AZ Humo sapiens.

15-0CT-2000. lu-APK-2000; 2000W0-US09688

05-APR-1999; 02-JUL-1999; 9905-0288950 9905-0346327

(CURI-) CORIXA CORP.

ƙeed SG, Xu J, Dillon DC;

WFI; 2000-638568/61.

A novel isolated polypeptide comprising an immunogenic portion of a treast cancer protein useful in the detection and treatment of breast caucer:

Claim 4; Page 77; 55pp; English.

The present sequence was isolated from a breast tumour cDNA library. It is provided in a specification relating to compounds for immunotherapy and diagnosts of breast cancer. Breast tumour antigens and the polynucleotities that encode them may be used in the production of a paramaceutical composition to be used in the treatment of breast cancer. Proliferated Teells and incubated antigen presenting cells are also required the polypeptides and polynucleotides may also be used to produce a vaccine.

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Sequence 1079 BP; 327 A; 229 C; 177 G; 346 T; 0 other;

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man polymacleotide SEQ ID NO 1200 CLNA; 1948 ů,

m.mar. nootropic; immanosuppressant; cytostatic; gene therapy; cancer, party.tarai nervous system; neuropathy; central nervous system; neuropathy; central nervous system; neuropath; claimer's; parkinson's disease; Huntington's disease; haemostatic; anjourophic lateral scherosis; shy-Drager Syndrome; chemotactic; c.e.mokinetic; thromolytic; drug screening; arthritis; inflammation; le.kamia; ss.

26-BEC-2000; 2000WG-0534263

21-JAH-2000; 2000US-0488725,
25-APR-2000; 2000US-0352317,
65-JUL-2000; 2000US-052931,
15-JUL-2000; 2000US-062931,
15-JUL-2000; 2000US-06291,
11-SEP-2000; 2000US-0654036,
11-SEP-2000; 2000US-0654036,
11-SEP-2000; 2000US-0654036,

2001-442253/47. EB: AAM39841. Asundi V. Chen R. Ma Y. Wehrman T. Xu C. Xue AJ. Goodrich R. Drmanac RT: Qian XB, Yang Y, Ren F, Wang D; Zhang J;

Claim 1; SEQ ID NO 1200; 10078pp; English.

for treating disorders

The invention relates to human nucleic acids (AM15795-AM161369) and the encoded polypeptides (AM36612-AM46131) with montropic, and incoming the polypeptides are useful in gase therapy. A composition containing a polypeptide or polynucleotide in gase therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system diseases, such as peripheral nervous injuries, peripheral nervopathy and incoming a polypeptide or polynucleotide of the invention may be used to treat diseases, such as extended the control solvenops. Activity, and incoming the activity and states, the state of the servous system suppression, and intrombolytic activity, cancer diagnosis and inflammation, leukeemias and each state of the activity and servous and inflammation, leukeemias and servous and activity and activity and theraptic activity. Kate: The sequence data for this patent did not form part of the printed specification.

Sequence 1946 BP; 614 A; 335 C; 436 G; 563 T; 0 other;

Query Match Best local Similarity 68.98; 69.98; Score y59; Db 22; Length 1948; Pred. No. 0;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1042
                                                                                                                         the invention relates to human nucleur acids (AAIS7798 AAI61169) and the encoded polypeptides (AAIS46211) with mootropic.

Inmunosuppressant and cytostatic activity. The polynuclectides are useful in gene therepy. A composition containing a polypeptide or polynuclectide containing a polynuclectide containing and period and containing and containing and containing and containing a polynuclectide containing and containing a polynuclectide containing a polynuclectide
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Tang YI, Wang J, Zhao QA,

Asundi V, Ch Wehrman T, X P, Goodrich R,

Chen R, Ma Y,
Xu C, Xue AJ,
R, Drmanac RT;

yang Y,

Ren F. Wang Zhang J:

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2001-442253/47. DB: AAM41627. Wang Z, Zhou P,

Claim 1; SEQ ID NO 4772; 10075pp; English such as central nervous system injuries -

ior

treating

disorders

Lery Match 88.5%; Score 95%; Di Sest Local Similarity 99.5%; Pred. No. 0; Matches 1079; Conservative 0; Mismatches

Sequence la56 bP;

613 A; 339 C; 442 G;

564 T; 0 other;

DB 22: Length 1958;

Indels

:-Gaps

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FLOUIT 4
FAISO783/C
AAISO783 Standard: CDMA:
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25-APR-2000;
05-JUL-2000;
15-JUL-2000;
03-AUG-2000;
14-SEP-2000;
19-CCT-2000;
25-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                      Human; noctropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; cNS; Altheimer's; Perkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thromtolytic; drug screening; arthritis; inflammation; buddemin; ss.
                                                                                                                                                                                                                                                                                                                                                                                      Humo sapiens.
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                                                                                                                                                                                                                                                  (HYSE-) HYSEQ INC
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2000US-0552317.
2000US-0558042.
2000US-0620312.
2000US-0623450.
2000US-06213450.
2000US-06213450.
2000US-0621345.
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933 actgccAntgfTTAGGTTTGCCCAAAACTCCCCAAACATCACAATGTTGTGGGTAAAC 874
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633 CTGGCCTTTTAATTICACTCTGATTICTICAACATTATACCTGTGAAATATCCTCTI 574
                                                                                                                                                           840 CATICACATGCACACATAAAGGTTTCTCATCATTTTGTACTTGGAAAAGGAATAATCT 639
                                                                                                                            780 cagcatgeetetggtgetteeaaatgagatgeataaggeaaageeaatteetgaca 639
753 CAGEATGECETTGGTGATTCTCCAAATGGATTCATAAGGCAAAAGCTCATTTCCTGACA 654
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(VLAA+) VLAAMS INIERUNIVERSITAIR INSE BIOTECHNOG Haylebroeck Dif:

Novel proteins used to treat inflammatory diseases, NF-kappaB related diseases and for improvement of anti-tumor treatments .

Claim 9; Fage 37-35; 46pp; English.

This sequence represents the gene encoding human CAO receptor associated protein (CRAP). CRAP is a functional protein capable of the tracting with the cytoplasmic domain of CAO and/or other receptors of the tumour necrosis factor (IRA) receptor superfamily such as CDO cand IRA receptor i, where the protein has no homology to TMA receptor associated factor (IRAP) proteins. The CO40 binding proteins can be used as modulators of the CD40 signalling pathway, especially to dispress and treat IRAF related, CD40 related, NF-Appab related and/or compared to the compared to fail tumour discusses. Diseases which may be treated include atherosclerosis, context is called the sclerosis, systemic lupus crythematosus, graft rejection, graft versus host disease, all for the improvement of anti-tumour treatments can be used to sensitive tumour cells to anti-tumour treatments and to screen for compounds which interfere with the proteins can be used to sensitive tumour cells to anti-tumour treatments and to screen for compounds which interfere with the proteins can be proteins with other protein components of the TRAF, CD40 or NF-Kappab related pathway.

Sequence 1920 BP; 559 A; 327 C; 435 G; 557 T; 2 other;

5 S Query Match Best Local Similarity 99.8%; Matches 1078; Conservative Score 508; DB 2 Pred. No. 0; 0; Mismatches БВ 21; Length 1920; 1; Indels 1; Gaps

Page 6

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RESULT 6
AAX28153/c
ID AAX28153 standard: DKA: 24% BP.
                                       AAX28153;
16-JUN-1999 (first entry)
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602 ettgeettittaattesetettgattiettesaeattatagstytgaaatacetteett 959
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782 CTSACTCIGGASCCICTTCCAILITETTTAAAACCANTIIIAACIGAIICAITCCITCCG 723
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812 CACCACATCGGTAACETCCGATCCCTTGATTTGTATCTCCTGCAATAAACTGTAG 783
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Best Local Similarity 99.8%;
Matches 1078; Conservative
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164 TAGJACATTATTCCTCCAAACAAACCTACCTTTCCAGAAGTGAAATTGTATTTGCA 1605
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121 typastccajctyjeogotataggascogitakaacicitacaaaactriacioccciiii 1765
               361
                                                                                                                                                                       ); score 908; DB 20; Length 2499;
); Pred. No. 0;
0; Mismatches 1; Indels 1; Gaps
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D)

Topulsomerase II binding protein; TopBP; anticancer agent; ds. Topulscrase II binding protein 1 coding sequence

JP11075856-A.

23-MAR-1599

17-SEP-1997; 97JP-0251544

17-SEP-1597: 97JP-0251544

WPI: 1999-257704/22. P-PSLB: AAY03182. (CHUS ) CHUGAI PHARM CO LTD. (ISUR/) ISURUO I.

120

TGGAATCCAGCTGCCAGCTGTAAGCACCGTTGAAAACTCTGACAGCTTTGTGCCCTTTT

ker Topoisomerase II: binding protein : useful as an anticancer agent

Disclosure; Fage 16-15; 28pp; Japanese.

This sequence encodes the topoiscomerase II Linding protein (TopBP) of the invention. The Topp protein is useful as an anticancer agent. TopBP can be used as the target molecule for anticancer agent.

Sequence 2499 BP; 720 A; 505 C; 567 G; 687 T; 0 other;

DB 20; Length 2499;

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RESULT

ACS8160/c

ACS8160/c

ACS8160/c

ACS8160/c

XX

ACS8160;

XX

Human; colon cancer antiyen nucle

XX

Human; colon cancer; colon cance

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                                 Colon cancer associated yene sequences, referred to as colon cancer antigens, useful for the treatment, prevention, and diagnosis of colon disorders such as colon cancer.
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CULTURALITICACICUGATUCII CONCACATIATAGCIGGAATHICCITCICATGA 542

665 CATCH991888CCTCC4931CCCCTL899TLT94TCTCCCTCCASAGATARTTT999CT9AC 784 HILLI H

601 505

AAC97991 to AAC98763 encode the human ocion carcor associated proteins, called human colon cancer antigens, given in AA69244 to AA694086. The Chiman colon cancer antigens can have cytostatic, cardioactive, muscular; netroprotective, influencedularcory, gysaecological, gastrointestinal, cultivatery, nephrotropic, antiniective and antibacterial activities, and can be used in gene therapy. The colon cancer antigen polynocleotides, and can be used in gene therapy. The colon cancer antigen polynocleotides, and creation and aliagnosis of colon disorders such as colon cancer. The proteins are useful for the prevention, cultratact, and diagnosis of colon disorders, such as colon cancer. The Colynocleotides may be used in dispositios and research, such as for color, cancer if ication, and as hybridisation probes. The proteins color cancer is entitled to prevent diseases such as incural disorders, immune color diseases such as incural disorders, such cardiors, lineartious diseases, and cardiovascular visorders, according to AA69872 and AA59407 represent sequences used in the exemplification of the present content of the present color and cancer is a color cancer. Claim 1; Fage 597; 2164pp; English.

Sequence 1296 bP; 376 A; 250 C; 326 G; 333 T; 11 other;

Ę \$ Ę, 5 Ę 8 Cb 3 usery Match 46.4 Best Local Similarity 59.5 Matches 771; Conservative 366 csjyjcasaccacacitgasaggatti.ccastatts.utctasgttgcacagaaga 425 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 426 coccequyaccactayaaatctaccacsgiccaytitticiaiccaggaagiccaaa 484
1061 CCCCAGTAATCACAAAACTACCAAGTCAGTTTTCTAATCCAAGAAGGTCAAA 1022 46.4%; Score 501; EB 21; Pred. No. 1.4e-246; 0; Mismatches 3; Length 1296; Gaps

545 ASITIACRASCASCASTANTOCAAGATTAGAGTICATIGGTATACCCAGGAGTAACTGG 902

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RESULG

AH15166

TD AAH15146 STALLITA; CDNA: loys EP

XX AAH15146;

XX AAH15146;

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XX AH15146;

XX Human cDNA sequence SEQ ID NO:13

XX Human; primer; setection: diagno

XX Human; primer; setection: diagno

XX Human; primer; setection: diagno

XX Primer sets for synthusizing pol

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                                                                                                                          Ota T, Isoyai I, Nishikawa I,
Ishii S, Sugiyama F, Wakamatsu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; primer; detection; diagnosis; untisense therap;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cDNA sequence SEQ ID NO:13205.
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Iuil-length obhas

Claim 8: Say ib 13205: 2537pp - CD ROM; anglish

An ine present invention describes primer sets for synthesising 5602 Cd [ill]-length chiks defined in the specification. Where a primer set comprises: (a) an object of primer and an object of complementary complementary strand of a polyvuclectide which complementary the 5602 handscalde sequences defined in the specification, where the collipcincolocials comprises at least 15 nucleotides or (t) a combination of an objective includes of the complementary to the complementary included the comprises at least 15 nucleotides or (t) a combination of a polyvucleotide which comprises a 5 fend of a polyvucleotide comprises a 5 fend of the combination of comprises a 5 fend of a polyvucleotide. The primer sets can be used in antisense therapy and comprises a 5 fend of the comprises therapy and comprise comprises a 5 fend of the comprise comprise comprise comprises a 5 fend of the comprise comprise comprises a 5 fend of the comprise comprise comprise comprises a 5 fend of the comprise comprise comprises a 5 fend of the comprise comprise comprises a 5 fend of the comprise comprises a 5 fend of the comprise comprises a 5 fend of the comprise comprise comprises a 5 fend of the comprise comprises a 5 fe

Sequence 1896 BP; 590 A; 327 C; 426 G; 555 I; 0 other;

Guery Match
Best Local Similarity 99.7
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74.18;

Score 795; DB 22; Length 1898; pred. No. u; u; Mismatches 2; Indels 1

2; indels l;

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ŗ 9 13 S 51 Ċ. 7 5 6 9 5 Ç E & E \$ 5 ξ. 1153 1253 1313 1373 1075 1133 ivis 111 4. 167 231 171 590 550 171 351 | CCCatalatactggcaatglttaggtttgccaaaaactccagacatccacaatgttg CARLARCERLARGICANA CONTROL CON 1 1 2 5 4 230 1154 505 554 5.25 1.7 410

Primer sets for synthesizing polymorlectides, particular, the full-length churs defined in the apectitication, and the first discharge and/or diagnosis of the abnormality of the proteins end and type for the proteins end type for the proteins end type for the proteins end to the proteins end type for the proteins e

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XX Human
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XX PR 29-JUL
CC 20-AAY
PR 11-JAN
PA (HELI-
XX WPI; 2
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                                                  The present invention describes primer sets for synthesising 5602 CC full-length cDNAs defined in the specification. Where a primer set CC full-length cDNAs defined in the specification. Where a primer set CC comprises: (a) an oligo-dT primer and an oligonucleotide complementary to the 5602 nucleotide sequences defined in the specification, where the CC cliponucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprising a sequence complementary to the CC complementary strand of a polynucleotide which comprises a 5'-end CC complementary to an oligonucleotide comprising a sequence complementary to a combination of CC complementary strand of a polynucleotide which comprises a 3'-end sequence complementary to a coligonucleotide which comprises a 3'-end sequence, where the coligonucleotide which comprises as 15 nucleotides and the combination of CC the 5'-end sequence/3'-end sequence is selected from those defined in CC the specification. The primers set useful for synthesising polynucleotides, and gene therapy. The primers are useful for synthesising polynucleotides, CC particularly full-length cDNAs. The primers are also useful for the CC particularly without any specialised methods. AAH03166 to AAH13628 and CC AAH3633 to AAH31872 represent human cDNA sequences; and AAH11629 to AAH13632 cof the present human amino acid sequences; and AAH13629 to AAH13632 cof the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer sets for synthesizing polynucleotides, particularly the 5502 full-length cDNAs defined in the specification, and for the detection and/or diagnosis of the abnormality of the proteins encoded by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID 4908;
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09-JUN-2000;
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2000JP-0183767.
2000JP-0241899.
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Otsuki
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Query Match

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RESULT 10
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DE Probe #5244 for gene expression in cervical cancer; ss.

KW Probe; human; microarray; gene KW Cervical cancer; ss.

CS Homo sapiens.

XX
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PD 09-AUG-2001; 2001W0-US00670.

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PN W0200157200; 2000US-0234687.

PR 30-JUN-2000; 2000US-0234687.

PR 30-JUN-2000; 2000US-0234687.

PR 30-AUG-2000; 2000US-0234687.

PR 30-AUG-2000; 2000US-033659.

PR 30-AUG-2000; 2000US-03366.

PR 30-AUG-2000; 2000US-0366.

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                                                                                                                                         The present invention relates to human single exon nucleic acid probes (SENP). The present sequence is one such probe. The SENPs are derived from human HeLa cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring luman gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer.
                                                               of diseases of the cervix, notably cervical cancer.
Note: The sequence data for this patent did not form specification, but was obtained in electronic format at ftp.wipo.int/puh/puhlishad nor communication.
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                                                                                                                                                                                                                                                                                                                                                      487pp;
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                                                                                                                                                                                                                                                                                                                                                                                                     ucleic acid probes useful cervical epithelial cells
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                                      The present invention relates to human single exon nucleic acid probes (SENP). The present sequence is one sprobe. The SENPs are derived from human HeLa cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1024
                                                                                                                                      Sequence 691 BP;
                                                                                                                                                                                                        expression, the probes are therefore useful in grading and/or of diseases of the cervix, notably cervical cancer. Note: The sequence data for this patent did not form part of useful increases, but was obtained in electronic formal directly specification, but was obtained in electronic formal directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 25; SEQ ID No 14417; 487pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               analyzing
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                                                                                                                                                                                   at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genome-derived single exon nucleic acid probes useful for zing gene expression in human cervical epithelial cells -
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2000US-0236359.
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                                  DB 22; |
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                                                                                           products. They can also be used to develop products for diagnosis and therapy. The proteins obtained may have cytckline activity, cell proliferation and differentiation activity, hasmatopoissis regulating activity, tissue growth regulating activity, reproductive hormone regulating activity, chemotractic/chemokinetic activity, hasmostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, tumour inhibition activity or other activities. The products can be used in forensic, gene therapy and chromosome mapping procedures The sequences can also be used for obtaining corresponding promoter sequences. The nucleic acids encoding the signal peptides can be used firecting extracellular secretion of a polypeptide or the insertion of a polypeptide into a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX40438 to AAX40715 represent 5' expressed sequence rags (ESTs) for human secreted proteins expressed in prostate, and encode the proteins given in AAX11716 to AAX11593 respectively. The proteins given represent the signal peptide and an N-terminal fragment of a secreted protein. The nucleic acid sequences can be used for producing secreted human gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human: secreted protein; EST; expressed sequence tag; diagnosis; forensic; gene therapy; chromosome mapping: slynal peptide; prostate; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopoiesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic; thrombolytic; anti-inflammatory; tumour inhibition; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 298; 675pp; English.
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PR 21-2

PR 21-2

PR 25-1

XX NOVE

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PR 25-1

XX NOVE

PR 25-1

AAZE

CC AA
                                                                                                                                                                                               AA265654 to AA269578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AA269579 to AA277440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; diagnosis; single nucleotide polymorphism; SNP; ds.
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23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-SEP-2001
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                                                                                                                                                                         effects from pharmaceutical agents acting
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                                                                                                                                                treatment
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                                                                                      . The SEQ ID NOS 2852, 2913, 3367, are not actually given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              agctaagtaggaacacaccc 1079
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 859;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9805-0082614
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                                                                         os 2852, 2913, 2974, 3035, actually given a sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2745pp;
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Best Local Similarity

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579

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other

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CC The present invention describes primer sets for synthesising 5602 CC comprises: (a) an oligo-dr primer and an oligocuclectide complementary to the complementary strand of a polynucleotide which complementary CC the 5602 nucleotide sequences defined in the specification, where the CC oligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprises as sequence complementary strand of a polynucleotide which comprises a 5'-end CC sequence and an oligonucleotide comprising a sequence, where the CC sequence and an oligonucleotide comprises a 3'-end sequence, where the CC oligonucleotide which comprises at least 15 nucleotides and the combination of CC the 5'-end sequence 3'-end sequence is selected from those defined in the specification. The primers sets can be used in antisense therapy and CC in gene therapy. The primers are useful for synthesising polynucleotides, particularly full length cDNAs. The primers are also useful for the detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs. The primers allow obtaining of the full-length CDNAs. The primers allow cottaining of the full-length CDNAs easily without any specialised methods. AAH03166 to AAH13628 and CC AAH3633 to AAH18742 represent human cDNA sequences; AAA92446 to CC AAB95893 represent human amino acid sequences; and AAH13629 to AAH13620 crepresent oligonucleotides, all of which are used in the exemplification of the present invention.
RESULT 1
AAAH1303
ID AAAH1
XX XX AAH1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer sets for synthesizing polynucleotides, particularly the 5602 full-length cDNAs defined in the specification, and for the detection and/or diagnosis of the abnormality of the proteins encoded by the
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02-MAY-2000;
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27-AUG-1595;
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Sugiyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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2000JF-0118776.
2000JF-0183767.
2000JP-0241899.
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A, Nagai K,
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Otsuki
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RESULT 15
AAA247119/c
ID AA24771
XX AAZ4771
XX Immune
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Query Match
Best Local Similarity
Matches 36; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunosuppressive; antiinflammatory; immunosuppressive; antiinflammatory; immunosuppressive; antiinflammatory; immunosuppressive; antiallergic; mouse; CD40 receptor associated protein; CRAP; cytoplasmic domain; tumour necrosis factor; TNF; receptor; superfamily; CD30; homology; TNF receptor associated factor; TRAF; modulator; signalling pathway; diagnosis; NF-KappaB; Jun; kinase; atherosclerosis; multiple sclerosis arthritis; systemic lupus erythematosus; graft rejection; allergy; graft versus host disease; autoimmune disease; ds.
                                                                                                                                                                                                                        of the tumour necrosis factor (TNF) receptor superfamily such as CD30 and TNF receptor I, where the protein has no homology to TNF receptor associated factor (TRAF)-proteins. The CD40 binding proteins can be used as modulators of the CD40 signalling pathway, especially to diagnose and treat TRAF-related, CD40-related, NF-kappaB related and/or Jun (kinase)-related diseases, and for the improvement of anti-tumour diseases. Diseases which may be treated include atheroscierosis, graft rejection, graft versus host disease, allergy, and autoimmune disease. The proteins can be used to sensitize tumour cells to anti-tumour treatments and to screen for compounds which interfere with the interaction of the proteins with other protein components of the TRAF CD40 or NF-kappaB related anatheav.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel proteins used to treat inflammatory diseases, NF-kappaB related diseases and for improvement of anti-tumor treatments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        associated protein (CRAP). CRAP is a functional protein capable of interacting with the cytoplasmic domain of CD40 and/or other receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P-PSDB; AAY56020
                                                                                                                                                       Sequence 1312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (VLAA-) VLAAMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse CD40 receptor associated protein gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA247119 standard; cDNA; 1312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W09955859-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence represents the gene encoding mouse CD40 receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-062029/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 41-43; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Remacle JEFJG,
                                                                                                                                                                                                           or NF-kappaB related
         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                       BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INTERUNIVERSITAIR INST BIOTECHNOG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98EP-0201392
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                                                                                                                                                       359
                              3.3%;
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                                                                                                                                                    279
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      0;
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   Score 36; DB Pred. No. 4.8
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Pred. No. 1.5e-12;
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                                                                                                                                                                                                        pathway.
                                                                                                                                                 352 G;
                                                                                                                                                    322 T;
                           DB 21; I
                                                                                                                                                 0 other;
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                                                          Length 1312;
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Indels

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Gaps

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149 agrittrictaatccaagaagg 169 agittictaatccaagaagg 479

459

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RESULT 1
AAH33140
  Matches
                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                  expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the provention, diagnosis and treatment of colorectal carcinomas and cancers. Assi7196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the proteins are collectively known as colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH3254: to AAH37195 and AAG73514 to AAG77788 represent human cancer associated nucleic acid molecules (N) and proteins (P).
                                                                                                                                                                     present invention.

N.B. Payes 666 to 682 and page 7053 of the missing at time of publication, meaning no SEQ ID 6.3:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 2344; 9803pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; colon cancer; colorectal carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-SEP-2001
                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding 4277 human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-235357/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-SEP-2000; 2000WO-US26524
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human colon cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH33140;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 417
Conservative
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                                                                                                                          BP; 118
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99US-0163280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antigen encoding cDNA SEQ ID NO:156
                                                                                                                          Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              colon cancer antigen;
; chromosome X; ss.
                     100.0%;
                                                                                                                        61 C;
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0;
                     Score 21;
Pred. No.
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                                                                                                                          65 G; 172
Mismatches
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                        1.5;
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                                                                                                                          1 other
                                                                                                                                                                                             sequence listing were sequences are present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis; detection;
                                             22; Length 117;
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Gaps
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RESULT 17
AAV55038
ID AAV550
AC AAV550
XX AAV550
XX Inhibi
KW Inhibi
KW Prolif
OS Homo s
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XX Homo s
XX Hopes
FT CDS
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FT CDS
FT TOJO
PN Hopes
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                                                                                                                                                                                                                                                                                                                              This sequence encodes the human XIAP protein, which is a inhibitor of C apoptosis protein (IAP), and can be used in the method of the invention. The method is for enhancing apoptosis in cells from a mammal with proliferative disease by treatment with a compound that inhibits compounds are used to treat proliferative diseases, specially cancers of compounds are used to treat proliferative diseases, specially cancers of covary, breast, pancreas, lymph nodes, skin, blood, lung, brain, kidney, criver nasopharnyx, thyroid,central nervous system, prostate, colon, crectum, cervix or endometrium, particularly to increase their sensitivity to chemotherapeutic agents of the IAP or NAIP proceins are detected in many cancers and are associated with poor prognosis, cresistance to chemotherapeutic agents and mutations in p53 (it is suggested that wild-type p53 suppresses transcription of the IAP or NAIP companies.) Transgenic animals are used for testing the effects of antisense collaponucleotides and for screening for the inhibitors.
                                                                                                                          Query Match
Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Baird S,
Tsang B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inducing apoptosis in proliferative mammalian cells with inhibitor of IAP or NAIP polypeptide - also methods for prognosis based on presence of IAP and NAIP, specifically applied to cancers involving
      2498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 13; Fig 1; 147pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         p53 mutations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV55038 standard; cDNA; 5232
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                                                                                                                                                                                                                                                                                   Sequence 5232 BP; 1579 A; 861 C; 1062 G; 1728 T; 2 other;
                                 459 agtttttctaatccaagaagg 479
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DB; AAW69294.
agtttttctaatccaagaagg 2518
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                                                                                                                              Conservative
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e disease;
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34..1527
/*tag= a
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/product= XIAP
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                                                                                                                                                     Score 21; pred. No.
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                                                                                                                              Mismatches
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                                                                                                                                                                                   Length 5232;
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AC AAC3
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DT 06-C
XX
DT 06-C
XX
Huma
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GE Huma
XX
GE 11-b
PR 26-1
PR 06-C
PR WPI
CC 10-C
PR OF
CC 11-C
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RESULT 19
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Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway;
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                                                                        Arabidopsis thaliana.
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114-OCT 1999
121-OCT 1999
121-OCT 1999
221-OCT 1999
221-OCT 1999
221-OCT 1999
221-OCT 1999
221-OCT 1999
225-OCT 1999
226-OCT 1999
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23-NOV-1998;
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                       21-APR-1999;
                                                                                     diagnosis;
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                                       28-OCT-1999
                                                                     Homo sapiens
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Pred. No. 4.8;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              treatment. N.B. The Sand 3367,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA265654 to AA269578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AA269579 to AA277440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the generic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         effects from pharmaceutical agents acting on a disease as well as trastmant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19
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                                                                                                                                                                                           WO200142467-A2
                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                      Cervical cancer;
                                                                                                                                                                                                                                                           Human cervical
                                                                                                                                                                                                                                                                                    19-SEP-2001
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                     Schlegel R,
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2001-375006/39
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                                           MILLENNIUM PREDICTIVE
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2000US-0220114.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                    Deeds J,
                                                                                                                                              2000WO-US33312
                                                                                                                         99US-0169681.
                                                                                                                                                                                                                                                             cancer marker nucleic acid
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                                                                                                                                                                                                                                      cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2745pp;
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Pred. No.
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                                                                                                                                                                                                                                      pre-malignant condition; gene therapy;
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                                           MEDICINE
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                     Zhao X;
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Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                     and
and
                                                                                                                                                                                                     progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for inhibiting cervical cancer in a patient. The nucleic acids may also be useful for gene therapy.
                                                                                                                                                                                                                                                                                                                                         Claim 1;
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                                                                       319 acagcaaacctacctttcc 337
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                                                                                                    Conservative
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RESULT 2
AAH71896
ID AAH7
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AX CETV
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2000US-0189315.
2000US-0203791.
2000US-0210600.
2000US-0220114.
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The invention relates to novel genes (AAH68727-AAH73383) associated with cervical cancer with cytostatic activity. The nucleic acids and encoded polypeptides are useful: to assess if a patient; is afflicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for

inhibiting cervical cancer in a patient. useful for gene therapy.

The nucleic acids may also

Claim 1;

Page 626; 1051pp; English.

isolated nucleic acid for diagnosing and treating the cancer -

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RESULT 2
AAH70689
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ID AAH70689
ID AAH7
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21-DEC-1999;
14-MAR-2000;
12-MAY-2000;
09-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                          Sequence
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acagcaaacctaccttcc
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19; Conser
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llarity 100.0%;
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                                                                                                                                                                                        128 A;
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                                                                                                                                                                                        106 C;
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Pred.
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                                                                                                                                                                                        114 G;
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                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy;
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AAH72662/c
ID AAH72662 standard; cDNA; 454 BP
XX
AC AAH72662;
XX
DT 19-SEP-2001 (first entry)
DE Human cervical cancer marker nu
XX
Cervical cancer; cytostatic; pr
XX Human cervical cancer; cytostatic; pr
XX Human cervical cancer; cytostatic; pr
XX Cervical cancer vith cytostatic
XX D8-DEC-1999; 99US-0171350.
PR 21-DEC-1999; 99US-0171310.
PR 21-DEC-1999; 99US-0171
Q
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                Human; secreted protein; gene therapy; protein therapy; cancer; weight; tumour; chromosome mapping; forensic; haematological disease; allergy; inflammation; cell proliferation; viral infection; wound healing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to novel genes (AAH68727-AAH73383) associated with cervical cancer with cytostatic activity. The nucleic actids and encoded polypeptides are useful; to assess if a patient is afflicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for inhibiting cervical cancer in a patient. The nucleic acids may also be
                                                                                                                                                      Human
                                                                                                                                                                                                          18-MAY-1999
                                                                                                                                                                                                                                                                                                             AAX22252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 787; 1051pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                         249
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                                                                                                                                                                                                                                                                                                                                                               26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             isolated nucleic acid for diagnosing for assessing and detecting compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                         ACAGCAAACCTACCTTTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acagcaaacctacctttcc 337
                                                                                                                                                 secreted protein gene 42 clone HSNAD72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
19; Conser
                                                                                                                                                                                                                                                                                                          standard; DNA;
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Matches

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Conservative

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Mismatches

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Gaps

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Score 19; Pred. No.

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Length 861;

Query Match Best Local Similarity

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The invention relates to nucleic acid sequences (AAX22211 to AAX22282)
CC encoding human secreted proteins (AAY0133 to AAY01454). The secreted compositing human secreted proteins (AAY0133) to AAY01454). The secreted compositing human sequences are deposited with the ATCC under deposit number atCC 209138, 209139 or 209141. Host cells containing vectors comprising the nucleic acid sequences are used for the recombinant expression of the nucleic acid sequences are useful for preventing, treating or ameliorating medical conditions e.g.
CC by protein or gene therapy. Pathological conditions can be also diaynosed by determining the amount of the new polynucleotides. The nucleic acid conditions to the new polynucleotides of the nucleic acid compared to the sequences, or its fragments, are useful for chromosome identification and amping; as antisonse and triplex-forming therapeutics; in gene therapy; for (forensic) identification of individuals; as molecular compared to the sequences or specific mRNA; in the preparation of oligomers and to raise anti-DNA antibodies. Antibodies are useful as immunoassay reagents (including for in vivo imaging) and conservation of the polynucleotide and complex for disorders may be treated with the polynucleotide and complex for disorders may be treated with the polynucleotide and complex for inflammation, cancer or other forms of cell proliferation, viral composition of embryonic stem cells, to modulate weight, appetite, behaviour etc. and as food additive or preservative. The presents sequence represents a done according a burgan secretary region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-JUL-1997
18-AUG-1997
18-AUG-1997
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22-JUL-1997;
22-JUL-1997;
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Greene JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 190-191; 251pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                binding
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P-PSDB; AAY01424.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acids encoding secreted human proteins - potentially useful for treating and diagnosing diseases and identifying specific
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16-JUL-1997;
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                                          sequence represents a gene encoding a human secreted protein
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Hu J,
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97US-0056359
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97US-0055985
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97US-0055726.
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Ni J, Ros
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                         for gene number and clone identification).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Young PE;
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861 BP;

222

A.

226 C;

164 G;

245 T;

4 other;

AAQ78892;

AAQ78892 standard; DNA; 1864

ВΡ

17-DEC-1995

(first entry)

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RESULT 2
AAQ78892
                                                                                                                                                  RESULT 1
AAN71405
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                                                                                                   Query Match
Best Local 9
                                                                                          Matches
                                                                                                                                                                   A DNA sequence coding for a heterologous polypeptide which can be expressed in and secreted from filamentous fungi is claimed. Pref the DNA sequence codes for bovine preprochymosin, M. meihei preprocarboxyl protease or A. niger preproglucoamylase. Also new are vectors consisting of the DNA sequence plus an operably-linked signal sequence. The vectors may also include a sequence which increases transformation efficiency, e.g. ANS-1.
                                             1161 trattttaaatattaacta 1179

    with secretion of prod. from
transformed fungi

                                                                                                                                                                                                                                                                 Example: Fig 13; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                           Cullen D,
                                                                                                                                                  Sequence 1864 BP;
                                                                                                                                                                                                                                                                                                               New DNA sequences tor
                                                                                                                                                                                                                                                                                                                                      WPI; 1987-095049/14
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29-AUG-1985;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Trichoderma; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Enzyme; fungal expression vector; Aspergillus expression vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence of ANS-1 which increases transformation efficiency.
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            28
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                                                                                           15; Conserv
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                                                                                                                                                                                                                                                                                                                                                         Gray GL,
                                                                                          Conservative
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85US-0771374
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                                                                                                                                                786 A; 210 C; 44 G;
                                                                                                     100.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                   ng polypeptide in filamentous fungi
the cells, and new vectors and
                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                            Lawlis
                                                                                                                                                732 T;
                                                                                                     DB 8;
15;
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                                                                                                             Length 1864;
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RESULT 29
AAF92529/C
ID AAF925
XX AAF925
XX 16-MAY
DT 16-MAY
DT 16-MAY
XX BAT T2
XX Human:
KW taste
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                                                                                                                                                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                               The sequence represents the A. nidulans ANS-1 sequence which is included in the construction of transformation vectors for recombinant protein expression and secretion from a filamentous fungus host. The sequence increases the transformation efficiency of the vector. This illustrates the main claims of the patent, which provide a vector containing (i).DNA encoding a heterologous polypeptide (chymosin, prochymosin, preprochymosin, Aspergillus niger glucoamylase, Humicola grisea glucoamylase or M. miehei carboxyl protease) and (ii) a secretory signal peptide, and a filamentous fungus (Aspergillus, Trichoderma, Neurospora, Podospora, Endothia, Mucor, Cochiobolus or Pyriclaria, especially A. nidulans, A. awamori or T. reesei) transformed with the vector for recombinant protein (enzyme) production.
                                                                                                                                                                                                                          1161
   15-MAR-2001
                                          Rattus
                                                                                                                                                                         AAF92529 standard;
                                                                                                                                                                                                                                                                                                               Sequence 1864 BP; 819 A; 193 C; 120 G; 732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vectors and DNA for expressing polypeptide(s) in filamentous funging include secretory signal sequences that are native or foreign to heterologous polypeptide(s), such as chymosin or glucoamylase.
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07-JUL-1986;
                        WO200118050-A2
                                                                                                             Rat T2R02 nucleotide sequence
                                                                                                                                   16-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
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                                                                                 taste
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vector; transformation;
                                                                                                                                                                                                                                           44 ttattttaaatattaacta 62
                                                                                                                                                                                                                                                                Local Similarity
les 19; Conser
                                                         ; rat; mouse; T2R; taste receptor; G-protein coupled receptor; transduction G-protein coupled receptor; identification; tong sensory neuron; taste cell; taste modulator; food; signalling pathway; ds.
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fig 13A-13B;
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                                                                                                                                 (first entry)
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86US-0882224
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RESULT 3
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probes for the identification of taste cells, as the nucleic acids are specifically expressed in taste cells. They also serve as tools for the generation of taste topographic maps that elucidate the relationship between the taste topographic maps that elucidate the relationship to taste centres in the brain. The taste modulators are useful for pharmacological and genetic modulation of taste signalling pathways. Modulatory compounds comprising T2R proteins can therefore be used in food and pharmaceutical industries to customise taste, for e.g. to decrease the bitter taste of food or drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF92502 to AAF92572 represent nucleic acids which encode taste transduction G-protein coupled receptors designated T2R proteins. AAB87731 to AAB87830 to AAB87731 to AAB87830 to Consensus sequences from the present invention. The T2R proteins are taste modulators. The nucleic acids are useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (REGC )
                                                                                                                                            chemokinetic; thrombolytic; drug screening; arthritis; leukaemia; ss.
                                                                                                                                                                                        peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral scierosis; Shy-Drager Syndrome; chemotactic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acids encoding the T2R family of G-protein coupled taste receptors, useful for identifying taste modulators that can be used food and pharmaceutical industries to customize taste, for e.g. to decrease the bitter taste of food.
                                              WO200153312-A1
                                                                                                                                                                                                                                                                                                                Human polynucleotide SEQ ID NO 1516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAI59313 standard; cDNA; 2706
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26-JUL-2001
                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                     Human;
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                                                                                                                                                                                                                                                           nootropic; immunosuppressant; cytostatic; gene therapy; cancer;
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19; Conserv
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US DEPT HEALTH
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SERVICES
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26-DEC-2000; 2000WO-US34263.

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RESULT 31
AAX13252;
ID AAX13252;
XX AXX13252;
XX 19-MAR-1999 (first of the content of the conten
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Matches 19
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03-AUG-2000;
14-SEP-2000;
19-OCT-2000;
29-NOV-2000;
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25-APR-2000;
09-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, Activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polypucleotides are useful in gene therapy. A composition containing a polypeptide or polypucleotide of the invention may be used to treat diseases of the peripheral nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tang
Wang
Zhao
                                                                                                                                                                                                                                Enterococcus faecalis; contig;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel nucleic acids and polypeptides, useful such as central nervous system injuries -
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Zhou P,
                                                                                                                                                                                                            attenuation; computer readable medium;
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2000US-0552317.
2000US-05598042.
2000US-0659812.
2000US-0653450.
2000US-0653450.
2000US-0653191.
2000US-0693036.
                                                                                                                                                                                                                                                                             faecalis genome contig SEQ ID NO:315
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Wehrman T, Xu C, Xue
Goodrich R, Drmanac
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Xue AJ,
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Yang Y,
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06-MAY-1997;
16-MAY-1997;
                                                                                                              28-SEP-2000; 2000WO-US26524
                                                                                                                                       05-APR-2001
                                                                                                                                                                                                                    colorectal carcinoma;
                                                                                                                                                                                                                             Human; colon
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                                                                                                                                                                                          sapiens
                                                                                                                                                                                                                                                     colon cancer antigen encoding cDNA SEQ ID NO:810.
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19; Conserv
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9703-0044031.
9703-0046655.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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0;

WPI; 2001-235357/24

Ruben SM,

Barash SC,

Birse

CE,

Rosen

CA;

(HUMA-) HUMAN GENOME SCI INC

29-SEP-1999; 03-NOV-1999;

99US-0157137 99US-0163280

P-PSDB; AAG74323

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RESULT 3
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps. by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB7789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                             Tango-77; human; II-1ra; cytokine superfamily; inflammation; inhibition; interleukin-1 receptor; II-1r, regulation; astma; rheumatoid arthritis; chronic myeloganous leukaemia; psoriasis; inflammatory bowel disease; growth factors; treatment; II-1 receptor complex; BAC; ss.
New isolated nucleic acid encoding the new used to inhibit inflammation and to screen
                                                    WPI; 1999-153692/13.
                                                                                                                                                    02-JUL-1998;
04-AUG-1997;
                                                                                                                                                                                                     03-AUG-1998;
                                                                                                                                                                                                                                                                    WO9906426-A1
                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                 Human IL-1ra BAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX02970;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX02970 standard; DNA; 1361 BP
                                                                                                                                                                                                                                     11-FEB-1999
                                                                                                                  (MILL-) MILLENNIUM BIOTHERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 291 BP; 86 A; 47 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B. Pages 666 to 682 and page 7053 of the sequence listing were
issing at time of publication, meaning no sequences are present
EQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ω
u
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gaaaacacaaccataaat 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gaaaacacaaccataaat 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cacids encoding 4277 human colon cancer-associated polypeptides, for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 2772;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                    98US-0091650
97US-0054646
                                                                                                                                                                                                   98WO-US16102
                                                                                                                                                                                                                                                                                                                                                                                                                contiguous DNA sequence 15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 G; 96 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 47;
              human cytokine Tango-77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 291;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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for specific modulators

P-PSDB; AAB41553 WPI; 2000-602362/57

RA,

Leach

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RESULT
AAC75762
Db
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                   31-MAR-1999;
02-APR-1999;
05-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                               vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant immunostimulant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; hypotensive; dermatological; immunosuppressive; antiinflammatory; antiviral; antibacterial; antiinqal; antirheumatic; antithyroid; antithanaxmic; gene therapy; cancer; proliferative disorder; hypertensic; neurodegenerative disorder; osteoarthritis; graft vs host disease; cardiovascular disease; diabetes mellitus; hypothyroidism; SCID; AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autoimmune disorder; asthma; allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound; bone damage; cartilage damage; antiinflammatory disease; coagulation; homes and complete the content of the combined immunodeficiency; malaria; autoimmune disorder; asthma; allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound; homes damage; cartilage damage; antiinflammatory disease; coagulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX0295,-x03048 and AAX22301-x22304 are overlapping BAC genomic sequences containing alternatively spliced forms of human IL-Ira. Such fragments are used in the method of the invention which describes the isolation of a novel human TANGO-77 encoding nucleic acid and protein. Tango-77 is a member of the cytokine superfamily that is expected to inhibit inflammation by binding to the interleukin-1 receptor (IL-IR). I may also bind to a new receptor so could regulate other cellular processes associated with acute or chronic inflammation, e.g. asthma, chronic myelogenous leukaemia, rheumatoid arthritis, psoriasis and inflammatory bowel disease. It may also induce or suppress interleukins, cytokinus and growth factors. Modulators of this protein are used to treat of prevent conditions associated with abnormal levels of inflammation, or activity of IL-1 or its receptor complex.
                                                                                                                                                                                                                                                                    05-OCT-2000
                                                                                                                                                                                                                               31-MAR-2000; 2000WO-US08621.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC75762
                                                                                           (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                            WO20005:173-A2
                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                      thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; open reading frame; ORFX; detection; cytostatic; hepatotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human OkfX ORF1317 polynucleotide sequence SEQ ID NO:2633.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1361 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             851 ttuaatattaactatttc 868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
nes [8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ttaaatattaactatttc 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Figure 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; cDNA; 1569
                                                                                                                                990S-0127636
990S-0127728
2000US-0540763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                        99US-0127607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       462 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 226pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       226 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     231 G; 442 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1361;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 35
AAQ22695/c
ID AAQ226
XX AAQ226
XX AAQ226
XX AAQ226
XX Sequen
XX Sequen
XX Carbon
XX Ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dρ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC antiporiatic; antiparkinsonian; notropic; neuroprotective; costeopathic; anticonvulsant; antiarthritic; immunosuppressant; cc immunostimulant; cardiant; thrombolytic; coagulant; vasotropic; cc antidiabetic; hypotensive; dermatological; immunosuppressive; cc antithyroid; and antianaemic. The sequences can be used for determining cc the presence of or predisposition to, or preventing or treating cc pathological conditions associated with an ORFX-associated disorder. The proteins and nucleic acids may be used to treat concers, cc proliferative disorders, neurodegenerative disorders, osteoarthritis, cc proliferative disorders, neurodegenerative disorders, osteoarthritis, cc praft vs host disease, cardiovascular disease, diabetes mellitus, cn hypertension, hypothyroidism, cholesterol ester storage, systemic lupus cerythematosus, severe combined immunodeficiency (SCID), AIDS, viral, callergies, aplastic anaemia, burns, wounds, bone and cartilage damage, coagulation; to inhibit thrombosis; and as a contraceptive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                            08-SEP-1990;
19-JUN-1991;
WPI; 1992-089407/12
                                                               Dworkin MB,
                                                                                                                (BOEH ) BOEHRINGER INGELHEI.
                                                                                                                                                                                                                                                                 08-SEP-1990;
                                                                                                                                                                                                                                                                                                                           12-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                     DE4028618-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Carbon metabolism; pyruvate formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence encoding mitochondrial NAD(P)+-dependent malate enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JUL-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ22695 standard; DNA; 1752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ22695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43357, which represent the human ORFX open reading frames 1 to 3161. The GRFX sequences have activities such as: cytostatic; hepatotropic; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel nucleic acids and peptides derived from open reading frame X, useful for treating e.g. cancers, proliferative disorders, neurodegenerative disorders and cardiovascular disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      839 acattcacatgcacacat 856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           540 acattcacatgcacacat 557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity hes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 1878–1879; 5507pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                         Leober G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      вP;
                                                                                                                                                                      90DE-4028618
91DE-4120178
                                                                                                                                                                                                                                                              90DE-4028618
                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
1..1752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   356 A; 487 C; 358 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                         Krystek
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 18;
Pred. No.
                                                      m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                   Maurer-Fogy I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   367 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB
49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 1569;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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RESULT 36
AAQ23258/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QΥ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SXCCCCCX PX P P X R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                    studying carbon metabolism in cells, a purificn. and assay
                                                                      WPI; 1992-114355/14.
P-PSDB; AAR22356.
                                                                                                                                            19-JUN-1591;
08-SEP-1590;
                                                                                                                                                                                                                                                                                                                                 CDS
                                                                                                   Dworkin MB,
                                                                                                                                                                                              19-MAR-1592
                                                                                                                                                                                                                                                          polyA_signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The inventors claim mitochondrial NAD(P)+-dependent malate enzyme and DNA encoding it. AAR21845 has 5' and 3' non-coding regions. The enzyme carachyses conversion of malate to pyruvate. Both the DNA and the enzyme are useful for studying carbon metabolism in rapidly dividing cells, esp. pyruvate formation from amino acids.
                                                                                                                        (BOEH ) BOEHRINGER INGELHEIM
                                                                                                                                                                           23-AUG-1991;
                                                                                                                                                                                                                   W09204448-A
                                                                                                                                                                                                                                                                              mat_peptide
                                                                                                                                                                                                                                                                                                   sig_peptide
                                                                                                                                                                                                                                                                                                                                                                                                       Mitochondrial
                                                                                                                                                                                                                                                                                                                                                                 Homo
                                                                                                                                                                                                                                                                                                                                                                               C-metabolism; tumour; pyruvate; T-lymphocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                            31-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ23258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ23258 standard; DNA; 1923 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1416 TGGAAAAATATAAACATT 1399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1752 BP; 562 A; 326 C; 386 G; 478 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Page 12-13; 20pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mitochondrial NAD(P)-dependent malate enzyme - used to study formation of pyruvate from aminoacid(s) in tumour cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB; AAR21845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.7%;
ilarity 100.0%;
Conservative
                                                                                                   Loeber G,
                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                    NAD(P)+-dependent malate enzyme
                                                                                                                                           91DE-4120178
90DE-4028618
                                                                                                                                                                          91WO-EP01602
                                                                                                                                                                                                                                                         /*tag= c
1905..1910
                                                                                                                                                                                                                                                                            /*tag= b
150..1844
                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
90..1844
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/note= "homologous to poly(A) signal AATAAA"
                                                                                                                                                                                                                                                                                                  /product= malate_enzyme
90..149
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                                                                                                   Krystek E,
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Pred. No.
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                                                                                                 Maureriog I,
                                   a and DNA encoding it - for also specific antibodies for
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Claim 1; Page 46 + Fig

3; 60pp; German

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RESULT 37
AAF18038/c
ID AAF180
XX AAF180
XX AAF180
XX AAF180
XX Human;
XW Cardio
XW Human;
XW Gastro
KW Gastro
CC Gassoci
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Best Local
Polynucleotide sequences AAF17982 - AAF18424 ercode human lung cancer associated proteins represented in AAB58106 - AAB58548. Lung cancer associated proteins and polynucleotide sequences, their agonists, and antagonists may have neuroprotective; cytostatic; cardioactive; immunomodulatory; muscular active general; vulnerary; gastrointestinal general; nephrotropic; antiinfective; gynecological; or antibacterial activity. The invention also includes antibodies specific for the protein or polynucleotide sequences. The lung cancer associated polynucleotide sequences may be used for detection of lung cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; runy consective; immunomodulatory; muscular active; various cardioactive; immunomodulatory; muscular active; gynecological; gastrointestinal; nephrotropic; antlinfective; gynecological; antibacterial; diagnosis; neural disorder; immune disorder; reantibacterial; diagnosis; neural disorder; immune disorder; disor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             basis for the design of oligonucleotides. These were used in PCR for amplification of malate enzyme encoding DNA in a CDNA bank prepd. from fibrosarcoma HS913. Amplified fragments were subcloned in pUC18, sequenced and used to probe the fibrosarcoma bank. A 1923bp insert was isolated and cloned in Bluescript KSr. The poly(A+) tail is not included in this sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1505
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antigens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-587514/55
P-PSDB; AAB58162.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ROSE/) HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; lung cancer associated protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      preparations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer associated gene sequences, referred gens, useful for treatment, prevention, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lung cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                       Page 536-537; 1425pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     associated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000WO-US05918
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0124270
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score
Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neuroprotective; cytostatic;
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of disorders
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RESULT 38
AAQ13822/c
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Matches 18
                The sequence was obtd. from 3 overlapping clones isolated from a human [24 cell line cDNA library. The DNA can be used to express the Gap b3 protein which is a transformation-dependent cell surface glycoprotein. The protein may be used to produce antibodies and these, or the DNA sequences, can be used to detect and quantify levels of Gap b3 protein or mRNA in biological samples. A high level of the protein is indicative of certain cancers. See also AAu13821-Q13824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gastrointestinal, pulmonary, cardiovascular, renal, and proliferative disorders. The proteins may also be used in the treatment of wounds are infectious diseases. Polynucleotide sequences AAF18425 - AAF18433 and peptide AAB58549 are used in the course of the invention for the identification and characterisation of the polynucleotide and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chromosome identification, as chromosome markers, and for diagnostic or research purposes. The proteins may be used disorders such as neural, immune, muscular, reproductive,
                                                                                                                                                                                                                                                                                                                                                                           CDS
                                                                                                                        Claim 1;
                                                                                                                                                         constructs
                                                                                                                                                                    DHA
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                                                                                                                                           sequences encoding galactoprotein b3 - produced structs also antibodies to Gap b3 used to detect ult in elevated expression of protein.
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RESULT 39
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Anti-integrin alpha3 KW
chemotherapeutic dru
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FH Key
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FT CDS
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          RESULT 40
AAA68247/c
ID AAA682
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AC AAA682
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DT 27-OCT
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Matches 18
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Best Local
                                                                                                                                                      27-OCT-2000
                                              AAA68247;
                                                                         AAA68247 standard;
                                                                                                                                                                                                                                                                                                                              This sequence encodes the human alpha3 integrin protein. The alpha3 integrin sequence is targeted by the anti-integrin alpha3 antibody of the invention. The anti-integrin alpha 3 antibody or its antigen binding fragment are for use as anti-tumour agents, and diagnostic reagent compositions. They can also be used in a chemotherapeutic drug.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-integrin alpha3 antibody and chemotherapeutic drug - anti-tumour agents and diagnostic reagent compositions
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18; Conserv
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                                                                         DNA; 41708
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AACB6106/C
ID AACB6106 standard; cf
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AC AACB6106;
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DT 29-AUG-2001 (first e
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DT Complete genome of ba
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Complete genome of ba
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DB Complete genome of ba
XX
DB Screening assay; ss.
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DS Bacteriophaye 77.
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PN WO200146383-A2.
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Best Local Similarity
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01-DEC-1995;
02-DEC-1995;
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03-JUN-1999;
28-SEP-1999;
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                                                                                                                                  Complete genome of bacteriophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 41708 BP; 15607
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-DEC-1999;
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                                                                                                                                                                  (first entry)
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99US-0168777.
99US-0454252.
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99US-0326144.
99US-0407804.
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                                                                                                                                                                                                                                  41708
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Bacteriophage: antimicrobial; genome; identification; antibacterial; bacterial growth inhibition; bacterial infection; ds.
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Identifying a bacteriophage coding region for treating bacterial infections comprises identifying a nucleic acid encoding a product inhibits bacteria when a bacteriophage infects a bacterium

Example 3; Page 141-151; 456pp; English.

The present invention describes a method for identifying a bacteriophage coding region encoding a product active on an essential bacterial target. The method comprises identifying a nucleic acid sequence encoding a gene product that provides a bacteria inhibiting function when an uncharacterised bacteriophage infects a pathogenic bacterium. The compound active on a target of a bacteriophage inhibitor protein in a bacteria is used to treat or prevent a bacterial infection in an animal. AAA68243 to AAA69442 and AAB16523 to AAB16524 represent bacteriophage nucleotide and protein sequences which are used the present invention. in the exemplification of.

5898 Ç 8088 G; 12115 T; 0 other;

1.7%; So larity 100.0%; E Conservative 0; Score 18; Pred. No. Mismatches DB 52; 21; 0; Length 41708; Indels 0, Gaps 0;

77

inhibitor; bacteriophage 77; ORF 104; phage 770RF104;

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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                               30283 CTTCATGACCTGTAATAA 30266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence represents the genome of Bacteriophage 77. The growth inhibitory gene product of ORF 104 interacts with DnaI derived from S. aureus, to form the basis of a screening assay. DnaI polypeptides and polynucleotides are useful for treating microbial, preferably bacterial, especially staphylococcal, infections, DnaI polypeptides and polynucleotides are useful for biological, ediagnostic, prophylactic, clinical and therapeutic use, and as components in databases useful for search analyses as well as in sequence analysis
                                                                                                                                                                                                            gene
New nucleic acid that is a 5' expressed sequence obtaining cDNAs and genomic DNAs that correspond
                                                                                                   26-FEB-1999;
                                                                                                                                              06-SEP-2000
                                                                                                                                                                   EP1033401-A2
                                                                                                                                                                                                                       Human;
                                                                                                                                                                                                                                                                  06-OCT-2000
                                                                                                                                                                                                                                                                                                          AAC31204 standard; cDNA; 212 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preferably
                                                                                                                        21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 41708 BP;
                                                                              (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                            957 cttcatgacctgtaataa 974
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12-OCT-2000; 2000US-0689952
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                                                                                                                                                                                                           therapy;
                                                        Milne Edwards
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bacterial, diseases
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                                                                                                                                                                                                          chromosome mapping;
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                                                                                                   99US-0122487
                                                                                                                                                                                                                      expressed
                                                                                                                                                                                                                                            protein 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             15607 A; 5898 C; 8088 G;
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                                                        Duclert A,
                                                                                                                                                                                                            mapping; ss.
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                                                                                                                                                                                                                                            SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                            ID NO:
                                                        Giordano
                                                                                                                                                                                                                   secreted protein; cDNA isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                     52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             12115 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                             Length 41708;
tag (5' EST) for to 5'ESTs and for
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Best Local
mRNAS encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAS of DAYA+ RNAS derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences
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                                                                                                                                                                                                                                                                                                                                             New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining CDNAs and genomic DNAs that correspond to 5'ESTs and for
                                                                                                                                                                                             The present sequence is one of a large number of 5' ESTs derived from
                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dumas Milne Edwards J.
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                                                                                                                                                                                                                                                    SEQ ID 14774; 71pp + CD-ROM; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chromosome mapping; ss.
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                                                                                                                                                                                                                                                                                                                    gene therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Duclert A,
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                                                                                                                                                                                                                                                                                                                 chromosome mapping procedures
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hes 0;
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RESULT 4
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21-DEC-1999;
14-MAR-2000;
12-MAY-2000;
09-JUN-2000;
21-JUL-2000;
                                                                                                                                               The invention relates to novel genes (AAH68727-AAH73383) associated with cervical cancer with cytostatic activity. The nucleic acids and encoded polypeptides are useful: to assess if a patient is afflicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for inhibiting cervical cancer in a patient. The nucleic acids may also be
                                                                                                                                                                                                                                                                                                                                      New
and
                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                              Schlegel R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              derived from the 5' ends of mRNAs and even in those cases where longer CDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                isolated nucleic acid for diagnosing and treating cervical cancer for assessing and detecting compounds for treating the cancer \cdot
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                                                                                                     329 BP; 117
                                                                                                                                                                                                                                                                                              Page 639; 1051pp; English
                                                                                                                                     gene therapy.
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2000US-0210600
2000US-0220114
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990S-0171350
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                          1.6%;
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         Score 17; DB Pred. No. 1.5
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                          DB 22; |
1.5e+02;
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thes 0;
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                                           Length 329;
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21-KAY-1999;
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01-JUN-1999;
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18-JUN-1999
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Search completed: April 30, 2002, 10:55:50 Job time: 11019 sec

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## ALIGNMENTS

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17-JUL-1998; 24-DEC-1997; 24-DEC-1997; 17-JUL-1998; Reed SG, (CORI-) CORIXA CORP 08-JUL-1999 Breast tumour breast cancer DNA encoding human breast tumour protein immunogenic fragment. W09933869-A2 22-DEC-1998; Homo sapiens û6-SEP-1999 AAX84210; AAX84210 standard; cDNA; 1001 BP X J; (first entry) protein; immunogenic fragment; development; therapy; ss. 98US-0118627. 97US-0998253. 97US-0998255. 98US-0118554. 98WO-US27416 vaccine; detection;

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New breast tumour protein genes used, in vaccines for immunotherapy, or for diaynosis of breast cancer

WPI; 1999-405486/34.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence encodes a human breast tumour protein immunogenic fragment of the invention. The polypeptides or nucleic acids encoding them are useful in vaccines and pharmaceutical compositions for manufacture of medicaments for inhibiting the development of breast cancer in a patient
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                                                 The present sequence was isolated from a breast tumour cDNA library. It is provided in a specification relating to compounds for immunotherapy and diagnosis of breast cancer. Breast tumour antigens and the polynucleotides that encode them may be used in the production of a pharmaceutical composition to be used in the treatment of breast cancer proliferated I cells and incubated antigen presenting cells are also required. The polypeptides and polynucleotides may also be used to produce a vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                                                     A novel isolated polypeptide comprising an immunogenic portion breast cancer protein useful in the detection and treatment of
                                                                                                                                                                                                                                                                                                Claim 4;
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02-JUL-1999;
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